

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property
Organization
International Bureau



(43) International Publication Date
19 May 2005 (19.05.2005)

PCT

(10) International Publication Number
WO 2005/044981 A2

(51) International Patent Classification⁷: C12N
(21) International Application Number: PCT/US2004/027403

(74) Agent: HILLMAN, Lisa, M., W.; McDonnell Boehnen
Hulbert and Berghoff LLP, 300 South Wacker Drive, Suite
3200, Chicago, IL 60606 (US).

(22) International Filing Date: 20 August 2004 (20.08.2004)

(81) Designated States (unless otherwise indicated, for every
kind of national protection available): AE, AG, AL, AM,
AT, AU, AZ, BA, BB, BG, BR, BW, BY, BZ, CA, CH, CN,
CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, EG, ES, FI,
GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, JP, KE,
KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD,
MG, MK, MN, MW, MX, MZ, NA, NI, NO, NZ, OM, PG,
PH, PL, PT, RO, RU, SC, SD, SE, SG, SK, SL, SY, TJ, TM,
TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, YU, ZA, ZM,
ZW.

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:
10/693,059 23 October 2003 (23.10.2003) US
10/720,448 24 November 2003 (24.11.2003) US
10/727,780 3 December 2003 (03.12.2003) US
10/757,803 14 January 2004 (14.01.2004) US
60/543,480 10 February 2004 (10.02.2004) US
10/780,447 13 February 2004 (13.02.2004) US
10/826,966 16 April 2004 (16.04.2004) US
PCT/US04/13456 30 April 2004 (30.04.2004) US
PCT/US04/16390 24 May 2004 (24.05.2004) US

(84) Designated States (unless otherwise indicated, for every
kind of regional protection available): ARIPO (BW, GH,
GM, KE, LS, MW, MZ, NA, SD, SL, SZ, TZ, UG, ZM,
ZW), Eurasian (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM),
European (AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI,
FR, GB, GR, HU, IE, IT, LU, MC, NL, PT, RO, SE, SI,
SK, TR), OAPI (BF, BJ, CF, CG, CI, CM, GA, GN, GQ,
GW, ML, MR, NE, SN, TD, TG).

(71) Applicant (for all designated States except US): SIRNA
THERAPEUTICS, INC. [US/US]; 2950 Wilderness
Place, Boulder, CO 80301 (US).

Published:

— without international search report and to be republished
upon receipt of that report

(72) Inventors; and

(75) Inventors/Applicants (for US only): HAEBERLI, Peter
[US/US]; 705 7th Street, Berthoud, CO 80513 (US). MC-
SWIGGEN, James [US/US]; 4866 Franklin Drive, Boul-
der, CO 80301 (US).

For two-letter codes and other abbreviations, refer to the "Guid-
ance Notes on Codes and Abbreviations" appearing at the begin-
ning of each regular issue of the PCT Gazette.

(54) Title: RNA INTERFERENCE MEDIATED INHIBITION OF GENE EXPRESSION USING SHORT INTERFERING NU-
CLEIC ACID (SINA)

(57) Abstract: This invention relates to compounds, compositions, and methods useful for modulating gene expression using short
interfering nucleic acid (siNA) molecules. In particular, the instant invention features small nucleic acid molecules, such as short
interfering nucleic acid (siNA), short interfering RNA (siRNA), double-stranded RNA (dsRNA), micro-RNA (miRNA), and short
hairpin RNA (shRNA) molecules and methods used to modulate the expression of genes, such as expressed pseudogenes associated
with the maintenance or development of diseases, disorders, traits, and conditions in a subject or organism.



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**RNA INTERFERENCE MEDIATED INHIBITION OF GENE EXPRESSION
USING SHORT INTERFERING NUCLEIC ACID (siNA)**

This application is a continuation-in-part of International Patent Application No. PCT/US04/16390, filed May 24, 2004, which is a continuation-in-part of U.S. Patent Application No. 10/826,966, filed April 16, 2004, which is continuation-in-part of U.S. Patent Application No. 10/757,803, filed January 14, 2004, which is a continuation-in-part of U.S. Patent Application No. 10/720,448, filed November 24, 2003, which is a continuation-in-part of U.S. Patent Application No. 10/693,059, filed October 23, 2003, which is a continuation-in-part of U.S. Patent Application No. 10/444,853, filed May 23, 2003, which is a continuation-in-part of International Patent Application No. PCT/US03/05346, filed February 20, 2003, and a continuation-in-part of International Patent Application No. PCT/US03/05028, filed February 20, 2003, both of which claim the benefit of U.S. Provisional Application No. 60/358,580 filed February 20, 2002, U.S. Provisional Application No. 60/363,124 filed March 11, 2002, U.S. Provisional Application No. 60/386,782 filed June 6, 2002, U.S. Provisional Application No. 60/406,784 filed August 29, 2002, U.S. Provisional Application No. 60/408,378 filed September 5, 2002, U.S. Provisional Application No. 60/409,293 filed September 9, 2002, and U.S. Provisional Application No. 60/440,129 filed January 15, 2003. This application is also a continuation-in-part of International Patent Application No. PCT/US04/13456, filed April 30, 2004, which is a continuation-in-part of U.S. Patent Application No. 10/780,447, filed February 13, 2004, which is a continuation-in-part of U.S. Patent Application No. 10/427,160, filed April 30, 2003, which is a continuation-in-part of International Patent Application No. PCT/US02/15876 filed May 17, 2002, which claims the benefit of U.S. Provisional Application No. 60/292,217, filed May 18, 2001, U.S. Provisional Application No. 60/362,016, filed March 6, 2002, U.S. Provisional Application No. 60/306,883, filed July 20, 2001, and U.S. Provisional Application No. 60/311,865, filed August 13, 2001. This application is also a continuation-in-part of U.S. Patent Application No. 10/727,780 filed December 3, 2003. This application also claims the benefit of U.S. Provisional Application No. 60/543,480, filed February 10, 2004. The instant application claims the benefit of all the listed applications, which are hereby incorporated by reference herein in their entireties, including the drawings.

Field Of The Invention

The present invention relates to compounds, compositions, and methods for the study, diagnosis, and treatment of traits, diseases and conditions that respond to the modulation of target gene expression and/or activity. The present invention is also directed to compounds, compositions, and methods relating to traits, diseases and conditions that respond to the modulation of expression and/or activity of genes involved in gene expression pathways or other cellular processes that mediate the maintenance or development of such traits, diseases and conditions. Specifically, the invention relates to small nucleic acid molecules, such as short interfering nucleic acid (siNA), short interfering RNA (siRNA), double-stranded RNA (dsRNA), micro-RNA (miRNA), and short hairpin RNA (shRNA) molecules capable of mediating RNA interference (RNAi) against target gene expression. Such small nucleic acid molecules are useful, for example, in providing compositions for treatment of traits, diseases and conditions that can respond to modulation of gene expression in a subject or organism.

Background Of The Invention

The following is a discussion of relevant art pertaining to RNAi. The discussion is provided only for understanding of the invention that follows. The summary is not an admission that any of the work described below is prior art to the claimed invention.

RNA interference refers to the process of sequence-specific post-transcriptional gene silencing in animals mediated by short interfering RNAs (siRNAs) (Zamore *et al.*, 2000, *Cell*, 101, 25-33; Fire *et al.*, 1998, *Nature*, 391, 806; Hamilton *et al.*, 1999, *Science*, 286, 950-951; Lin *et al.*, 1999, *Nature*, 402, 128-129; Sharp, 1999, *Genes & Dev.*, 13:139-141; and Strauss, 1999, *Science*, 286, 886). The corresponding process in plants (Heifetz *et al.*, International PCT Publication No. WO 99/61631) is commonly referred to as post-transcriptional gene silencing or RNA silencing and is also referred to as quelling in fungi. The process of post-transcriptional gene silencing is thought to be an evolutionarily-conserved cellular defense mechanism used to prevent the expression of foreign genes and is commonly shared by diverse flora and phyla (Fire *et al.*, 1999, *Trends Genet.*, 15, 358). Such protection from foreign gene expression may have evolved in response to the production of double-stranded RNAs (dsRNAs) derived from

viral infection or from the random integration of transposon elements into a host genome via a cellular response that specifically destroys homologous single-stranded RNA or viral genomic RNA. The presence of dsRNA in cells triggers the RNAi response through a mechanism that has yet to be fully characterized. This mechanism appears to be different from other known mechanisms involving double stranded RNA-specific ribonucleases, such as the interferon response that results from dsRNA-mediated activation of protein kinase PKR and 2',5'-oligoadenylate synthetase resulting in non-specific cleavage of mRNA by ribonuclease L (see for example US Patent Nos. 6,107,094; 5,898,031; Clemens *et al.*, 1997, *J. Interferon & Cytokine Res.*, 17, 503-524; Adah *et al.*, 2001, *Curr. Med. Chem.*, 8, 1189).

The presence of long dsRNAs in cells stimulates the activity of a ribonuclease III enzyme referred to as dicer (Bass, 2000, *Cell*, 101, 235; Zamore *et al.*, 2000, *Cell*, 101, 25-33; Hammond *et al.*, 2000, *Nature*, 404, 293). Dicer is involved in the processing of the dsRNA into short pieces of dsRNA known as short interfering RNAs (siRNAs) (Zamore *et al.*, 2000, *Cell*, 101, 25-33; Bass, 2000, *Cell*, 101, 235; Bernstein *et al.*, 2001, *Nature*, 409, 363). Short interfering RNAs derived from dicer activity are typically about 21 to about 23 nucleotides in length and comprise about 19 base pair duplexes (Zamore *et al.*, 2000, *Cell*, 101, 25-33; Elbashir *et al.*, 2001, *Genes Dev.*, 15, 188). Dicer has also been implicated in the excision of 21- and 22-nucleotide small temporal RNAs (stRNAs) from precursor RNA of conserved structure that are implicated in translational control (Hutvagner *et al.*, 2001, *Science*, 293, 834). The RNAi response also features an endonuclease complex, commonly referred to as an RNA-induced silencing complex (RISC), which mediates cleavage of single-stranded RNA having sequence complementary to the antisense strand of the siRNA duplex. Cleavage of the target RNA takes place in the middle of the region complementary to the antisense strand of the siRNA duplex (Elbashir *et al.*, 2001, *Genes Dev.*, 15, 188).

RNAi has been studied in a variety of systems. Fire *et al.*, 1998, *Nature*, 391, 806, were the first to observe RNAi in *C. elegans*. Bahramian and Zarbl, 1999, *Molecular and Cellular Biology*, 19, 274-283 and Wianny and Goetz, 1999, *Nature Cell Biol.*, 2, 70, describe RNAi mediated by dsRNA in mammalian systems. Hammond *et al.*, 2000, *Nature*, 404, 293, describe RNAi in *Drosophila* cells transfected with dsRNA. Elbashir

et al., 2001, *Nature*, 411, 494 and Tuschl *et al.*, International PCT Publication No. WO 01/75164, describe RNAi induced by introduction of duplexes of synthetic 21-nucleotide RNAs in cultured mammalian cells including human embryonic kidney and HeLa cells. Recent work in *Drosophila* embryonic lysates (Elbashir *et al.*, 2001, *EMBO J.*, 20, 6877 and Tuschl *et al.*, International PCT Publication No. WO 01/75164) has revealed certain requirements for siRNA length, structure, chemical composition, and sequence that are essential to mediate efficient RNAi activity. These studies have shown that 21-nucleotide siRNA duplexes are most active when containing 3'-terminal dinucleotide overhangs. Furthermore, complete substitution of one or both siRNA strands with 2'-deoxy (2'-H) or 2'-O-methyl nucleotides abolishes RNAi activity, whereas substitution of the 3'-terminal siRNA overhang nucleotides with 2'-deoxy nucleotides (2'-H) was shown to be tolerated. Single mismatch sequences in the center of the siRNA duplex were also shown to abolish RNAi activity. In addition, these studies also indicate that the position of the cleavage site in the target RNA is defined by the 5'-end of the siRNA guide sequence rather than the 3'-end of the guide sequence (Elbashir *et al.*, 2001, *EMBO J.*, 20, 6877). Other studies have indicated that a 5'-phosphate on the target-complementary strand of a siRNA duplex is required for siRNA activity and that ATP is utilized to maintain the 5'-phosphate moiety on the siRNA (Nykanen *et al.*, 2001, *Cell*, 107, 309).

Studies have shown that replacing the 3'-terminal nucleotide overhanging segments of a 21-mer siRNA duplex having two-nucleotide 3'-overhangs with deoxyribonucleotides does not have an adverse effect on RNAi activity. Replacing up to four nucleotides on each end of the siRNA with deoxyribonucleotides has been reported to be well tolerated, whereas complete substitution with deoxyribonucleotides results in no RNAi activity (Elbashir *et al.*, 2001, *EMBO J.*, 20, 6877 and Tuschl *et al.*, International PCT Publication No. WO 01/75164). In addition, Elbashir *et al.*, *supra*, also report that substitution of siRNA with 2'-O-methyl nucleotides completely abolishes RNAi activity. Li *et al.*, International PCT Publication No. WO 00/44914, and Beach *et al.*, International PCT Publication No. WO 01/68836 preliminarily suggest that siRNA may include modifications to either the phosphate-sugar backbone or the nucleoside to include at least one of a nitrogen or sulfur heteroatom, however, neither application postulates to what extent such modifications would be tolerated in siRNA molecules, nor provides any further guidance or examples of such modified siRNA. Kreutzer *et al.*,

Canadian Patent Application No. 2,359,180, also describe certain chemical modifications for use in dsRNA constructs in order to counteract activation of double-stranded RNA-dependent protein kinase PKR, specifically 2'-amino or 2'-O-methyl nucleotides, and nucleotides containing a 2'-O or 4'-C methylene bridge. However, Kreutzer *et al.* similarly fails to provide examples or guidance as to what extent these modifications would be tolerated in dsRNA molecules.

Parrish *et al.*, 2000, *Molecular Cell*, 6, 1077-1087, tested certain chemical modifications targeting the unc-22 gene in *C. elegans* using long (>25 nt) siRNA transcripts. The authors describe the introduction of thiophosphate residues into these siRNA transcripts by incorporating thiophosphate nucleotide analogs with T7 and T3 RNA polymerase and observed that RNAs with two phosphorothioate modified bases also had substantial decreases in effectiveness as RNAi. Further, Parrish *et al.* reported that phosphorothioate modification of more than two residues greatly destabilized the RNAs *in vitro* such that interference activities could not be assayed. *Id.* at 1081. The authors also tested certain modifications at the 2'-position of the nucleotide sugar in the long siRNA transcripts and found that substituting deoxynucleotides for ribonucleotides produced a substantial decrease in interference activity, especially in the case of Uridine to Thymidine and/or Cytidine to deoxy-Cytidine substitutions. *Id.* In addition, the authors tested certain base modifications, including substituting, in sense and antisense strands of the siRNA, 4-thiouracil, 5-bromouracil, 5-iodouracil, and 3-(aminoallyl)uracil for uracil, and inosine for guanosine. Whereas 4-thiouracil and 5-bromouracil substitution appeared to be tolerated, Parrish reported that inosine produced a substantial decrease in interference activity when incorporated in either strand. Parrish also reported that incorporation of 5-iodouracil and 3-(aminoallyl)uracil in the antisense strand resulted in a substantial decrease in RNAi activity as well.

The use of longer dsRNA has been described. For example, Beach *et al.*, International PCT Publication No. WO 01/68836, describes specific methods for attenuating gene expression using endogenously-derived dsRNA. Tuschl *et al.*, International PCT Publication No. WO 01/75164, describe a *Drosophila in vitro* RNAi system and the use of specific siRNA molecules for certain functional genomic and certain therapeutic applications; although Tuschl, 2001, *Chem. Biochem.*, 2, 239-245,

- doubts that RNAi can be used to cure genetic diseases or viral infection due to the danger of activating interferon response. Li *et al.*, International PCT Publication No. WO 00/44914, describe the use of specific long (141 bp-488 bp) enzymatically synthesized or vector expressed dsRNAs for attenuating the expression of certain target genes.
- 5 Zernicka-Goetz *et al.*, International PCT Publication No. WO 01/36646, describe certain methods for inhibiting the expression of particular genes in mammalian cells using certain long (550 bp-714 bp), enzymatically synthesized or vector expressed dsRNA molecules. Fire *et al.*, International PCT Publication No. WO 99/32619, describe particular methods for introducing certain long dsRNA molecules into cells for use in
- 10 inhibiting gene expression in nematodes. Plaetinck *et al.*, International PCT Publication No. WO 00/01846, describe certain methods for identifying specific genes responsible for conferring a particular phenotype in a cell using specific long dsRNA molecules. Mello *et al.*, International PCT Publication No. WO 01/29058, describe the identification of specific genes involved in dsRNA-mediated RNAi. Pachuck *et al.*, International PCT
- 15 Publication No. WO 00/63364, describe certain long (at least 200 nucleotide) dsRNA constructs. Deschamps Depaillette *et al.*, International PCT Publication No. WO 99/07409, describe specific compositions consisting of particular dsRNA molecules combined with certain anti-viral agents. Waterhouse *et al.*, International PCT Publication No. 99/53050 and 1998, *PNAS*, 95, 13959-13964, describe certain methods
- 20 for decreasing the phenotypic expression of a nucleic acid in plant cells using certain dsRNAs. Driscoll *et al.*, International PCT Publication No. WO 01/49844, describe specific DNA expression constructs for use in facilitating gene silencing in targeted organisms.

- Others have reported on various RNAi and gene-silencing systems. For example,
- 25 Parrish *et al.*, 2000, *Molecular Cell*, 6, 1077-1087, describe specific chemically-modified dsRNA constructs targeting the unc-22 gene of *C. elegans*. Grossniklaus, International PCT Publication No. WO 01/38551, describes certain methods for regulating polycomb gene expression in plants using certain dsRNAs. Churikov *et al.*, International PCT Publication No. WO 01/42443, describe certain methods for modifying genetic
- 30 characteristics of an organism using certain dsRNAs. Cogoni *et al.*, International PCT Publication No. WO 01/53475, describe certain methods for isolating a Neurospora silencing gene and uses thereof. Reed *et al.*, International PCT Publication No. WO

- 01/68836, describe certain methods for gene silencing in plants. Honer *et al.*, International PCT Publication No. WO 01/70944, describe certain methods of drug screening using transgenic nematodes as Parkinson's Disease models using certain dsRNAs. Deak *et al.*, International PCT Publication No. WO 01/72774, describe certain
- 5 *Drosophila*-derived gene products that may be related to RNAi in *Drosophila*. Arndt *et al.*, International PCT Publication No. WO 01/92513 describe certain methods for mediating gene suppression by using factors that enhance RNAi. Tuschl *et al.*, International PCT Publication No. WO 02/44321, describe certain synthetic siRNA constructs. Pachuk *et al.*, International PCT Publication No. WO 00/63364, and
- 10 Satishchandran *et al.*, International PCT Publication No. WO 01/04313, describe certain methods and compositions for inhibiting the function of certain polynucleotide sequences using certain long (over 250 bp), vector expressed dsRNAs. Echeverri *et al.*, International PCT Publication No. WO 02/38805, describe certain *C. elegans* genes identified via RNAi. Kreutzer *et al.*, International PCT Publications Nos. WO
- 15 02/055692, WO 02/055693, and EP 1144623 B1 describes certain methods for inhibiting gene expression using dsRNA. Graham *et al.*, International PCT Publications Nos. WO 99/49029 and WO 01/70949, and AU 4037501 describe certain vector expressed siRNA molecules. Fire *et al.*, US 6,506,559, describe certain methods for inhibiting gene expression in vitro using certain long dsRNA (299 bp-1033 bp) constructs that mediate
- 20 RNAi. Martinez *et al.*, 2002, *Cell*, 110, 563-574, describe certain single stranded siRNA constructs, including certain 5'-phosphorylated single stranded siRNAs that mediate RNA interference in Hela cells. Harborth *et al.*, 2003, *Antisense & Nucleic Acid Drug Development*, 13, 83-105, describe certain chemically and structurally modified siRNA molecules. Chiu and Rana, 2003, *RNA*, 9, 1034-1048, describe certain chemically and
- 25 structurally modified siRNA molecules. Woolf *et al.*, International PCT Publication Nos. WO 03/064626 and WO 03/064625 describe certain chemically modified dsRNA constructs.

SUMMARY OF THE INVENTION

- This invention relates to compounds, compositions, and methods useful for
- 30 modulating gene expression using short interfering nucleic acid (siNA) molecules. This invention also relates to compounds, compositions, and methods useful for modulating

the expression and activity of other genes involved in pathways of gene expression and/or activity by RNA interference (RNAi) using small nucleic acid molecules. In particular, the instant invention features small nucleic acid molecules, such as short interfering nucleic acid (siNA), short interfering RNA (siRNA), double-stranded RNA (dsRNA), micro-RNA (miRNA), and short hairpin RNA (shRNA) molecules and methods used to modulate the expression of genes, including gene targets having RNA transcripts referred to by Genbank Accession Numbers shown in **Table I**.

A siNA of the invention can be unmodified or chemically-modified. A siNA of the instant invention can be chemically synthesized, expressed from a vector or enzymatically synthesized. The instant invention also features various chemically-modified synthetic short interfering nucleic acid (siNA) molecules capable of modulating target gene expression or activity in cells by RNA interference (RNAi). The use of chemically-modified siNA improves various properties of native siNA molecules through increased resistance to nuclease degradation *in vivo* and/or through improved cellular uptake. Further, contrary to earlier published studies, siNA having multiple chemical modifications retains its RNAi activity. The siNA molecules of the instant invention provide useful reagents and methods for a variety of therapeutic, veterinary, diagnostic, target validation, genomic discovery, genetic engineering, and pharmacogenomic applications.

In one embodiment, the invention features one or more siNA molecules and methods that independently or in combination modulate the expression of target genes encoding proteins, such as proteins that are associated with the maintenance and/or development of diseases, traits, disorders, and/or conditions as described herein or otherwise known in the art, such as genes encoding sequences comprising those sequences referred to by GenBank Accession Nos. shown in **Table I**, referred to herein generally as "target". The description below of the various aspects and embodiments of the invention is provided with reference to exemplary target genes referred to herein as gene targets. However, the various aspects and embodiments are also directed to other genes, such as gene homologs, transcript variants, and polymorphisms (e.g., single nucleotide polymorphism, (SNPs)) associated with certain genes. As such, the various aspects and embodiments are also directed to other genes that are involved in disease,

trait, condition, or disorder related pathways of signal transduction or gene expression that are involved, for example, in the maintenance or development of diseases, traits, conditions, or disorders described herein. These additional genes can be analyzed for target sites using the methods described for exemplary genes herein. Thus, the modulation of other genes and the effects of such modulation of the other genes can be performed, determined, and measured as described herein.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of a target gene or that directs cleavage of a target RNA, wherein said siNA molecule comprises about 15 to about 28 base pairs.

In one embodiment, the invention features a double stranded short interfering nucleic acid (siNA) molecule that directs cleavage of a target RNA via RNA interference (RNAi), wherein the double stranded siNA molecule comprises a first and a second strand, each strand of the siNA molecule is about 18 to about 28 nucleotides in length, the first strand of the siNA molecule comprises nucleotide sequence having sufficient complementarity to the target RNA for the siNA molecule to direct cleavage of the target RNA via RNA interference, and the second strand of said siNA molecule comprises nucleotide sequence that is complementary to the first strand.

In one embodiment, the invention features a double stranded short interfering nucleic acid (siNA) molecule that directs cleavage of a target RNA via RNA interference (RNAi), wherein the double stranded siNA molecule comprises a first and a second strand, each strand of the siNA molecule is about 18 to about 23 nucleotides in length, the first strand of the siNA molecule comprises nucleotide sequence having sufficient complementarity to the target RNA for the siNA molecule to direct cleavage of the target RNA via RNA interference, and the second strand of said siNA molecule comprises nucleotide sequence that is complementary to the first strand.

In one embodiment, the invention features a chemically synthesized double stranded short interfering nucleic acid (siNA) molecule that directs cleavage of a target RNA via RNA interference (RNAi), wherein each strand of the siNA molecule is about 18 to about 28 nucleotides in length; and one strand of the siNA molecule comprises

nucleotide sequence having sufficient complementarity to the target RNA for the siNA molecule to direct cleavage of the target RNA via RNA interference.

In one embodiment, the invention features a chemically synthesized double stranded short interfering nucleic acid (siNA) molecule that directs cleavage of a target
5 RNA via RNA interference (RNAi), wherein each strand of the siNA molecule is about 18 to about 23 nucleotides in length; and one strand of the siNA molecule comprises nucleotide sequence having sufficient complementarity to the target RNA for the siNA molecule to direct cleavage of the target RNA via RNA interference.

In one embodiment, the invention features a siNA molecule that down-regulates
10 expression of a target gene or that directs cleavage of a target RNA, for example, wherein the gene comprises protein encoding sequence. In one embodiment, the invention features a siNA molecule that down-regulates expression of a target gene or that directs cleavage of a target RNA, for example, wherein the gene comprises non-coding sequence or encodes sequence of regulatory elements involved in gene expression
15 (e.g., non-coding RNA).

In one embodiment, a siNA of the invention is used to inhibit the expression of target genes or a target gene family, wherein the genes or gene family sequences share sequence homology. Such homologous sequences can be identified as is known in the art, for example using sequence alignments. siNA molecules can be designed to target
20 such homologous sequences, for example using perfectly complementary sequences or by incorporating non-canonical base pairs, for example mismatches and/or wobble base pairs, that can provide additional target sequences. In instances where mismatches are identified, non-canonical base pairs (for example, mismatches and/or wobble bases) can be used to generate siNA molecules that target more than one gene sequence. In a non-limiting example, non-canonical base pairs such as UU and CC base pairs are used to generate siNA molecules that are capable of targeting sequences for differing targets that share sequence homology. As such, one advantage of using siNAs of the invention is that a single siNA can be designed to include nucleic acid sequence that is complementary to the nucleotide sequence that is conserved between the homologous
30 genes. In this approach, a single siNA can be used to inhibit expression of more than one gene instead of using more than one siNA molecule to target the different genes.

In one embodiment, a target RNA of the invention is an expressed pseudogene (see for example pseudogene sequences referred to by Genbank Accession Numbers in **Table I**). As used herein the term "disease related expressed pseudogene" refers to any expressed pseudogene that is associated with a disease, disorder, condition, or trait.

5 In one embodiment, the invention features a siNA molecule having RNAi activity against target RNA (e.g., coding or non-coding RNA), wherein the siNA molecule comprises a sequence complementary to any RNA sequence, such as those sequences having GenBank Accession Nos. shown in **Table I**. In another embodiment, the invention features a siNA molecule having RNAi activity against target RNA, wherein
10 the siNA molecule comprises a sequence complementary to an RNA having variant encoding sequence, for example other mutant genes not shown in **Table I** but known in the art to be associated with the maintenance and/or development of diseases, traits, disorders, and/or conditions described herein or otherwise known in the art. Chemical modifications as shown in **Table II** or otherwise described herein can be applied to any
15 siNA construct of the invention. In another embodiment, a siNA molecule of the invention includes a nucleotide sequence that can interact with nucleotide sequence of a target gene and thereby mediate silencing of gene expression, for example, wherein the siNA mediates regulation of gene expression by cellular processes that modulate the chromatin structure or methylation patterns of the gene and prevent transcription of the
20 gene.

In one embodiment, siNA molecules of the invention are used to down regulate or inhibit the expression of proteins arising from haplotype polymorphisms that are associated with a disease or condition. Analysis of genes, or protein or RNA levels can be used to identify subjects with such polymorphisms or those subjects who are at risk of
25 developing traits, conditions, or diseases described herein. These subjects are amenable to treatment, for example, treatment with siNA molecules of the invention and any other composition useful in treating diseases related to gene expression. As such, analysis of protein or RNA levels can be used to determine treatment type and the course of therapy in treating a subject. Monitoring of protein or RNA levels can be used to predict
30 treatment outcome and to determine the efficacy of compounds and compositions that

modulate the level and/or activity of certain proteins associated with a trait, disorder, condition, or disease.

In one embodiment of the invention a siNA molecule comprises an antisense strand comprising a nucleotide sequence that is complementary to a target polynucleotide sequence or a portion thereof. The siNA further comprises a sense strand, wherein said sense strand comprises a nucleotide sequence of a target polynucleotide sequence or a portion thereof, (e.g., about 15 to about 25 or more, or about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, or 25 or more contiguous nucleotides in a target polynucleotide sequence). In one embodiment, the target polynucleotide sequence is a target DNA. In one embodiment, the target polynucleotide sequence is a target RNA.

In one embodiment, the invention features a siNA molecule comprising a first sequence, for example, the antisense sequence of the siNA construct, complementary to a sequence or portion of sequence comprising sequence represented by GenBank Accession Nos. shown in **Table I**, and a second sequence, for example a sense sequence, that is complementary to the antisense sequence. Chemical modifications in **Table II** and described herein can be applied to any siNA construct (e.g., sense or antisense sequence) of the invention.

In one embodiment of the invention a siNA molecule comprises an antisense strand having about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides, wherein the antisense strand is complementary to a target RNA sequence or a portion thereof, and wherein said siNA further comprises a sense strand having about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides, and wherein said sense strand and said antisense strand are distinct nucleotide sequences where at least about 15 nucleotides in each strand are complementary to the other strand.

In another embodiment of the invention a siNA molecule of the invention comprises an antisense region having about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides, wherein the antisense region is complementary to a target DNA sequence, and wherein said siNA further comprises a sense region having about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23,

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14, 15, 16, 17, 18, 19, 20, 21, or 22 nucleotides, wherein said sense region and said antisense region are comprised in a linear molecule where the sense region comprises at least about 15 nucleotides that are complementary to the antisense region.

5 In one embodiment, a siNA molecule of the invention has RNAi activity that modulates expression of RNA encoded by one or more genes. Because various genes can share some degree of sequence homology with each other, siNA molecules can be designed to target a class of genes or alternately specific genes (e.g., polymorphic variants) by selecting sequences that are either shared amongst different gene targets or alternatively that are unique for a specific gene target. Therefore, in one embodiment, 10 the siNA molecule can be designed to target conserved regions of target RNA sequences having homology among several gene variants so as to target a class of genes with one siNA molecule. Accordingly, in one embodiment, the siNA molecule of the invention modulates the expression of one or both gene alleles in a subject. In another embodiment, the siNA molecule can be designed to target a sequence that is unique to a 15 specific target RNA sequence (e.g., a single allele or single nucleotide polymorphism (SNP)) due to the high degree of specificity that the siNA molecule requires to mediate RNAi activity.

In one embodiment, nucleic acid molecules of the invention that act as mediators of the RNA interference gene silencing response are double-stranded nucleic acid 20 molecules. In another embodiment, the siNA molecules of the invention consist of duplex nucleic acid molecules containing about 15 to about 30 base pairs between oligonucleotides comprising about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides. In yet another embodiment, siNA molecules of the invention comprise duplex nucleic acid molecules with overhanging 25 ends of about 1 to about 3 (e.g., about 1, 2, or 3) nucleotides, for example, about 21-nucleotide duplexes with about 19 base pairs and 3'-terminal mononucleotide, dinucleotide, or trinucleotide overhangs. In yet another embodiment, siNA molecules of the invention comprise duplex nucleic acid molecules with blunt ends, where both ends are blunt, or alternatively, where one of the ends is blunt.

30 In one embodiment, the invention features one or more chemically-modified siNA constructs having specificity for a target polynucleotide (e.g., RNA or DNA), such as

5 DNA encoding RNA sequences referred to herein by Genbank Accession number or such RNA sequences referred to herein by Genbank Accession number. In one embodiment, the invention features a RNA based siNA molecule (e.g., a siNA comprising 2'-OH nucleotides) having specificity for target polynucleotides (e.g., RNA or DNA) that includes one or more chemical modifications described herein. Non-limiting examples of such chemical modifications include without limitation phosphorothioate internucleotide linkages, 2'-deoxyribonucleotides, 2'-O-methyl ribonucleotides, 2'-deoxy-2'-fluoro ribonucleotides, "universal base" nucleotides, "acyclic" nucleotides, 5-C-methyl nucleotides, and terminal glyceryl and/or inverted deoxy abasic residue incorporation. These chemical modifications, when used in various siNA constructs, (e.g., RNA based siNA constructs), are shown to preserve RNAi activity in cells while at the same time, dramatically increasing the serum stability of these compounds. Furthermore, contrary to the data published by Parrish *et al.*, *supra*, applicant demonstrates that multiple (greater than one) phosphorothioate substitutions are well-tolerated and confer substantial increases in serum stability for modified siNA constructs.

In one embodiment, a siNA molecule of the invention comprises modified nucleotides while maintaining the ability to mediate RNAi. The modified nucleotides can be used to improve *in vitro* or *in vivo* characteristics such as stability, activity, and/or bioavailability. For example, a siNA molecule of the invention can comprise modified nucleotides as a percentage of the total number of nucleotides present in the siNA molecule. As such, a siNA molecule of the invention can generally comprise about 5% to about 100% modified nucleotides (e.g., about 5%, 10%, 15%, 20%, 25%, 30%, 35%, 40%, 45%, 50%, 55%, 60%, 65%, 70%, 75%, 80%, 85%, 90%, 95% or 100% modified nucleotides). The actual percentage of modified nucleotides present in a given siNA molecule will depend on the total number of nucleotides present in the siNA. If the siNA molecule is single stranded, the percent modification can be based upon the total number of nucleotides present in the single stranded siNA molecules. Likewise, if the siNA molecule is double stranded, the percent modification can be based upon the total number of nucleotides present in the sense strand, antisense strand, or both the sense and antisense strands.

One aspect of the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of a target gene or that directs cleavage of a target RNA. In one embodiment, the double stranded siNA molecule comprises one or more chemical modifications and each strand of the double-stranded siNA is about 21 nucleotides long. In one embodiment, the double-stranded siNA molecule does not contain any ribonucleotides. In another embodiment, the double-stranded siNA molecule comprises one or more ribonucleotides. In one embodiment, each strand of the double-stranded siNA molecule independently comprises about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides, wherein each strand comprises about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides that are complementary to the nucleotides of the other strand. In one embodiment, one of the strands of the double-stranded siNA molecule comprises a nucleotide sequence that is complementary to a nucleotide sequence or a portion thereof of the gene, and the second strand of the double-stranded siNA molecule comprises a nucleotide sequence substantially similar to the nucleotide sequence of the gene or a portion thereof.

In another embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of a target gene or that directs cleavage of a target RNA, comprising an antisense region, wherein the antisense region comprises a nucleotide sequence that is complementary to a nucleotide sequence of the gene or a portion thereof, and a sense region, wherein the sense region comprises a nucleotide sequence substantially similar to the nucleotide sequence of the gene or a portion thereof. In one embodiment, the antisense region and the sense region independently comprise about 15 to about 30 (e.g. about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides, wherein the antisense region comprises about 15 to about 30 (e.g. about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides that are complementary to nucleotides of the sense region.

In another embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of a target gene or that directs cleavage of a target RNA, comprising a sense region and an antisense region, wherein the antisense region comprises a nucleotide sequence that is complementary to a

nucleotide sequence of RNA encoded by the gene or a portion thereof and the sense region comprises a nucleotide sequence that is complementary to the antisense region.

In one embodiment, a siNA molecule of the invention comprises blunt ends, i.e., ends that do not include any overhanging nucleotides. For example, a siNA molecule comprising modifications described herein (e.g., comprising nucleotides having
5 Formulae I-VII or siNA constructs comprising "Stab 00"-"Stab 32" (Table II) or any combination thereof (see Table II)) and/or any length described herein can comprise blunt ends or ends with no overhanging nucleotides.

In one embodiment, any siNA molecule of the invention can comprise one or more
10 blunt ends, i.e. where a blunt end does not have any overhanging nucleotides. In one embodiment, the blunt ended siNA molecule has a number of base pairs equal to the number of nucleotides present in each strand of the siNA molecule. In another embodiment, the siNA molecule comprises one blunt end, for example wherein the 5'-end of the antisense strand and the 3'-end of the sense strand do not have any
15 overhanging nucleotides. In another example, the siNA molecule comprises one blunt end, for example wherein the 3'-end of the antisense strand and the 5'-end of the sense strand do not have any overhanging nucleotides. In another example, a siNA molecule comprises two blunt ends, for example wherein the 3'-end of the antisense strand and the 5'-end of the sense strand as well as the 5'-end of the antisense strand and 3'-end of the
20 sense strand do not have any overhanging nucleotides. A blunt ended siNA molecule can comprise, for example, from about 15 to about 30 nucleotides (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30 nucleotides). Other nucleotides present in a blunt ended siNA molecule can comprise, for example, mismatches, bulges, loops, or wobble base pairs to modulate the activity of the siNA molecule to mediate
25 RNA interference.

By "blunt ends" is meant symmetric termini or termini of a double stranded siNA molecule having no overhanging nucleotides. The two strands of a double stranded siNA molecule align with each other without over-hanging nucleotides at the termini. For example, a blunt ended siNA construct comprises terminal nucleotides that are
30 complementary between the sense and antisense regions of the siNA molecule.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of a target gene or that directs cleavage of a target RNA, wherein the siNA molecule is assembled from two separate oligonucleotide fragments wherein one fragment comprises the sense region and
5 the second fragment comprises the antisense region of the siNA molecule. The sense region can be connected to the antisense region via a linker molecule, such as a polynucleotide linker or a non-nucleotide linker.

In one embodiment, the invention features double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of a target gene or that
10 directs cleavage of a target RNA, wherein the siNA molecule comprises about 15 to about 30 (e.g. about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) base pairs, and wherein each strand of the siNA molecule comprises one or more chemical modifications. In another embodiment, one of the strands of the double-stranded siNA molecule comprises a nucleotide sequence that is complementary to a nucleotide
15 sequence of a gene or a portion thereof, and the second strand of the double-stranded siNA molecule comprises a nucleotide sequence substantially similar to the nucleotide sequence or a portion thereof of the gene. In another embodiment, one of the strands of the double-stranded siNA molecule comprises a nucleotide sequence that is complementary to a nucleotide sequence of a gene or portion thereof, and the second
20 strand of the double-stranded siNA molecule comprises a nucleotide sequence substantially similar to the nucleotide sequence or portion thereof of the gene. In another embodiment, each strand of the siNA molecule comprises about 15 to about 30 (e.g. about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides, and each strand comprises at least about 15 to about 30 (e.g. about 15, 16, 17, 18, 19, 20, 21,
25 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides that are complementary to the nucleotides of the other strand. The gene can comprise, for example, a gene that encodes sequences referred to in Table I.

In one embodiment, a siNA molecule of the invention comprises no ribonucleotides. In another embodiment, a siNA molecule of the invention comprises
30 ribonucleotides.

In one embodiment, a siNA molecule of the invention comprises an antisense region comprising a nucleotide sequence that is complementary to a nucleotide sequence of a target gene or a portion thereof, and the siNA further comprises a sense region comprising a nucleotide sequence substantially similar to the nucleotide sequence of the target gene or a portion thereof. In another embodiment, the antisense region and the sense region each comprise about 15 to about 30 (e.g. about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides and the antisense region comprises at least about 15 to about 30 (e.g. about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides that are complementary to nucleotides of the sense region. The target gene can comprise, for example, sequence encoding sequences referred to in Table I. In another embodiment, the siNA is a double stranded nucleic acid molecule, where each of the two strands of the siNA molecule independently comprise about 15 to about 40 (e.g. about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 23, 33, 34, 35, 36, 37, 38, 39, or 40) nucleotides, and where one of the strands of the siNA molecule comprises at least about 15 (e.g. about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24 or 25 or more) nucleotides that are complementary to the nucleic acid sequence of the gene or a portion thereof.

In one embodiment, a siNA molecule of the invention comprises a sense region and an antisense region, wherein the antisense region comprises a nucleotide sequence that is complementary to a nucleotide sequence of RNA encoded by a target gene, or a portion thereof, and the sense region comprises a nucleotide sequence that is complementary to the antisense region. In one embodiment, the siNA molecule is assembled from two separate oligonucleotide fragments, wherein one fragment comprises the sense region and the second fragment comprises the antisense region of the siNA molecule. In another embodiment, the sense region is connected to the antisense region via a linker molecule. In another embodiment, the sense region is connected to the antisense region via a linker molecule, such as a nucleotide or non-nucleotide linker. The target gene can comprise, for example, sequence encoding sequences referred to in Table I.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of a target gene or that

directs cleavage of a target RNA comprising a sense region and an antisense region, wherein the antisense region comprises a nucleotide sequence that is complementary to a nucleotide sequence of RNA encoded by the target gene or a portion thereof and the sense region comprises a nucleotide sequence that is complementary to the antisense region, and wherein the siNA molecule has one or more modified pyrimidine and/or purine nucleotides. In one embodiment, the pyrimidine nucleotides in the sense region are 2'-O-methyl pyrimidine nucleotides or 2'-deoxy-2'-fluoro pyrimidine nucleotides and the purine nucleotides present in the sense region are 2'-deoxy purine nucleotides. In another embodiment, the pyrimidine nucleotides in the sense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides and the purine nucleotides present in the sense region are 2'-O-methyl purine nucleotides. In another embodiment, the pyrimidine nucleotides in the sense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides and the purine nucleotides present in the sense region are 2'-deoxy purine nucleotides. In one embodiment, the pyrimidine nucleotides in the antisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides and the purine nucleotides present in the antisense region are 2'-O-methyl or 2'-deoxy purine nucleotides. In another embodiment of any of the above-described siNA molecules, any nucleotides present in a non-complementary region of the sense strand (e.g. overhang region) are 2'-deoxy nucleotides.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of a target gene or that directs cleavage of a target RNA, wherein the siNA molecule is assembled from two separate oligonucleotide fragments wherein one fragment comprises the sense region and the second fragment comprises the antisense region of the siNA molecule, and wherein the fragment comprising the sense region includes a terminal cap moiety at the 5'-end, the 3'-end, or both of the 5' and 3' ends of the fragment. In one embodiment, the terminal cap moiety is an inverted deoxy abasic moiety or glyceryl moiety. In one embodiment, each of the two fragments of the siNA molecule independently comprise about 15 to about 30 (e.g. about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides. In another embodiment, each of the two fragments of the siNA molecule independently comprise about 15 to about 40 (e.g. about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 23, 33, 34, 35, 36, 37, 38, 39, or 40) nucleotides. In a

non-limiting example, each of the two fragments of the siNA molecule comprise about 21 nucleotides.

- In one embodiment, the invention features a siNA molecule comprising at least one modified nucleotide, wherein the modified nucleotide is a 2'-deoxy-2'-fluoro nucleotide.
- 5 The siNA can be, for example, about 15 to about 40 nucleotides in length. In one embodiment, all pyrimidine nucleotides present in the siNA are 2'-deoxy-2'-fluoro pyrimidine nucleotides. In one embodiment, the modified nucleotides in the siNA include at least one 2'-deoxy-2'-fluoro cytidine or 2'-deoxy-2'-fluoro uridine nucleotide. In another embodiment, the modified nucleotides in the siNA include at least one 2'-
- 10 fluoro cytidine and at least one 2'-deoxy-2'-fluoro uridine nucleotides. In one embodiment, all uridine nucleotides present in the siNA are 2'-deoxy-2'-fluoro uridine nucleotides. In one embodiment, all cytidine nucleotides present in the siNA are 2'-deoxy-2'-fluoro cytidine nucleotides. In one embodiment, all adenosine nucleotides present in the siNA are 2'-deoxy-2'-fluoro adenosine nucleotides. In one embodiment,
- 15 all guanosine nucleotides present in the siNA are 2'-deoxy-2'-fluoro guanosine nucleotides. The siNA can further comprise at least one modified internucleotidic linkage, such as phosphorothioate linkage. In one embodiment, the 2'-deoxy-2'-fluoronucleotides are present at specifically selected locations in the siNA that are sensitive to cleavage by ribonucleases, such as locations having pyrimidine nucleotides.
- 20 In one embodiment, the invention features a method of increasing the stability of a siNA molecule against cleavage by ribonucleases comprising introducing at least one modified nucleotide into the siNA molecule, wherein the modified nucleotide is a 2'-deoxy-2'-fluoro nucleotide. In one embodiment, all pyrimidine nucleotides present in the siNA are 2'-deoxy-2'-fluoro pyrimidine nucleotides. In one embodiment, the
- 25 modified nucleotides in the siNA include at least one 2'-deoxy-2'-fluoro cytidine or 2'-deoxy-2'-fluoro uridine nucleotide. In another embodiment, the modified nucleotides in the siNA include at least one 2'-fluoro cytidine and at least one 2'-deoxy-2'-fluoro uridine nucleotides. In one embodiment, all uridine nucleotides present in the siNA are 2'-deoxy-2'-fluoro uridine nucleotides. In one embodiment, all cytidine nucleotides
- 30 present in the siNA are 2'-deoxy-2'-fluoro cytidine nucleotides. In one embodiment, all adenosine nucleotides present in the siNA are 2'-deoxy-2'-fluoro adenosine nucleotides.

In one embodiment, all guanosine nucleotides present in the siNA are 2'-deoxy-2'-fluoro guanosine nucleotides. The siNA can further comprise at least one modified internucleotidic linkage, such as phosphorothioate linkage. In one embodiment, the 2'-deoxy-2'-fluoronucleotides are present at specifically selected locations in the siNA that are sensitive to cleavage by ribonucleases, such as locations having pyrimidine nucleotides.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of a target gene or that directs cleavage of a target RNA comprising a sense region and an antisense region, wherein the antisense region comprises a nucleotide sequence that is complementary to a nucleotide sequence of RNA encoded by the gene or a portion thereof and the sense region comprises a nucleotide sequence that is complementary to the antisense region, and wherein the purine nucleotides present in the antisense region comprise 2'-deoxy-purine nucleotides. In an alternative embodiment, the purine nucleotides present in the antisense region comprise 2'-O-methyl purine nucleotides. In either of the above embodiments, the antisense region can comprise a phosphorothioate internucleotide linkage at the 3' end of the antisense region. Alternatively, in either of the above embodiments, the antisense region can comprise a glyceryl modification at the 3' end of the antisense region. In another embodiment of any of the above-described siNA molecules, any nucleotides present in a non-complementary region of the antisense strand (e.g. overhang region) are 2'-deoxy nucleotides.

In one embodiment, the antisense region of a siNA molecule of the invention comprises sequence complementary to a portion of a target polynucleotide sequence having sequence unique to a particular disease related allele, such as sequence comprising a single nucleotide polymorphism (SNP) associated with the disease specific allele. As such, the antisense region of a siNA molecule of the invention can comprise sequence complementary to sequences that are unique to a particular allele to provide specificity in mediating selective RNAi against the disease, condition, or trait related allele.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of a target gene or that

directs cleavage of a target RNA, wherein the siNA molecule is assembled from two separate oligonucleotide fragments wherein one fragment comprises the sense region and the second fragment comprises the antisense region of the siNA molecule. In another embodiment, the siNA molecule is a double stranded nucleic acid molecule, where each strand is about 21 nucleotides long and where about 19 nucleotides of each fragment of the siNA molecule are base-paired to the complementary nucleotides of the other fragment of the siNA molecule, wherein at least two 3' terminal nucleotides of each fragment of the siNA molecule are not base-paired to the nucleotides of the other fragment of the siNA molecule. In another embodiment, the siNA molecule is a double stranded nucleic acid molecule, where each strand is about 19 nucleotide long and where the nucleotides of each fragment of the siNA molecule are base-paired to the complementary nucleotides of the other fragment of the siNA molecule to form at least about 15 (e.g., 15, 16, 17, 18, or 19) base pairs, wherein one or both ends of the siNA molecule are blunt ends. In one embodiment, each of the two 3' terminal nucleotides of each fragment of the siNA molecule is a 2'-deoxy-pyrimidine nucleotide, such as a 2'-deoxy-thymidine. In another embodiment, all nucleotides of each fragment of the siNA molecule are base-paired to the complementary nucleotides of the other fragment of the siNA molecule. In another embodiment, the siNA molecule is a double stranded nucleic acid molecule of about 19 to about 25 base pairs having a sense region and an antisense region, where about 19 nucleotides of the antisense region are base-paired to the nucleotide sequence or a portion thereof of the RNA encoded by the target gene. In another embodiment, about 21 nucleotides of the antisense region are base-paired to the nucleotide sequence or a portion thereof of the RNA encoded by the target gene. In any of the above embodiments, the 5'-end of the fragment comprising said antisense region can optionally include a phosphate group.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that inhibits the expression of a target RNA sequence (e.g., wherein said target RNA sequence is encoded by a gene involved in a pathway of gene expression), wherein the siNA molecule does not contain any ribonucleotides and wherein each strand of the double-stranded siNA molecule is about 15 to about 30 nucleotides. In one embodiment, the siNA molecule is 21 nucleotides in length. Examples of non-ribonucleotide containing siNA constructs are combinations of

stabilization chemistries shown in **Table II** in any combination of Sense/Antisense chemistries, such as Stab 7/8, Stab 7/11, Stab 8/8, Stab 18/8, Stab 18/11, Stab 12/13, Stab 7/13, Stab 18/13, Stab 7/19, Stab 8/19, Stab 18/19, Stab 7/20, Stab 8/20, Stab 18/20, Stab 7/32, Stab 8/32, or Stab 18/32 (e.g., any siNA having Stab 7, 8, 11, 12, 13, 14, 15, 17, 18, 19, 20, or 32 sense or antisense strands or any combination thereof).

In one embodiment, the invention features a chemically synthesized double stranded RNA molecule that directs cleavage of a target RNA via RNA interference, wherein each strand of said RNA molecule is about 15 to about 30 nucleotides in length; one strand of the RNA molecule comprises nucleotide sequence having sufficient complementarity to the target RNA for the RNA molecule to direct cleavage of the target RNA via RNA interference; and wherein at least one strand of the RNA molecule optionally comprises one or more chemically modified nucleotides described herein, such as without limitation deoxynucleotides, 2'-O-methyl nucleotides, 2'-deoxy-2'-fluoro nucleotides, 2'-O-methoxyethyl nucleotides etc.

In one embodiment, a target RNA of the invention comprises sequence encoding a protein.

In one embodiment, target RNA of the invention comprises non-coding RNA sequence (e.g., miRNA, snRNA siRNA etc.).

In one embodiment, the invention features a medicament comprising a siNA molecule of the invention.

In one embodiment, the invention features an active ingredient comprising a siNA molecule of the invention.

In one embodiment, the invention features the use of a double-stranded short interfering nucleic acid (siNA) molecule to inhibit, down-regulate, or reduce expression of a gene or that directs cleavage of a target RNA, wherein the siNA molecule comprises one or more chemical modifications and each strand of the double-stranded siNA is independently about 15 to about 30 or more (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29 or 30 or more) nucleotides long. In one embodiment, the siNA molecule of the invention is a double stranded nucleic acid molecule comprising one or

more chemical modifications, where each of the two fragments of the siNA molecule independently comprise about 15 to about 40 (e.g. about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 23, 33, 34, 35, 36, 37, 38, 39, or 40) nucleotides and where one of the strands comprises at least 15 nucleotides that are complementary to

5 nucleotide sequence of target RNA or a portion thereof. In a non-limiting example, each of the two fragments of the siNA molecule comprise about 21 nucleotides. In another embodiment, the siNA molecule is a double stranded nucleic acid molecule comprising one or more chemical modifications, where each strand is about 21 nucleotide long and where about 19 nucleotides of each fragment of the siNA molecule are base-paired to the

10 complementary nucleotides of the other fragment of the siNA molecule, wherein at least two 3' terminal nucleotides of each fragment of the siNA molecule are not base-paired to the nucleotides of the other fragment of the siNA molecule. In another embodiment, the siNA molecule is a double stranded nucleic acid molecule comprising one or more chemical modifications, where each strand is about 19 nucleotide long and where the

15 nucleotides of each fragment of the siNA molecule are base-paired to the complementary nucleotides of the other fragment of the siNA molecule to form at least about 15 (e.g., 15, 16, 17, 18, or 19) base pairs, wherein one or both ends of the siNA molecule are blunt ends. In one embodiment, each of the two 3' terminal nucleotides of each fragment of the siNA molecule is a 2'-deoxy-pyrimidine nucleotide, such as a 2'-deoxy-thymidine.

20 In another embodiment, all nucleotides of each fragment of the siNA molecule are base-paired to the complementary nucleotides of the other fragment of the siNA molecule. In another embodiment, the siNA molecule is a double stranded nucleic acid molecule of about 19 to about 25 base pairs having a sense region and an antisense region and comprising one or more chemical modifications, where about 19 nucleotides of the

25 antisense region are base-paired to the nucleotide sequence or a portion thereof of the RNA encoded by the target gene. In another embodiment, about 21 nucleotides of the antisense region are base-paired to the nucleotide sequence or a portion thereof of the RNA encoded by the target gene. In any of the above embodiments, the 5'-end of the fragment comprising said antisense region can optionally include a phosphate group.

30 In one embodiment, the invention features the use of a double-stranded short interfering nucleic acid (siNA) molecule that inhibits, down-regulates, or reduces expression of a target gene or that directs cleavage of a target RNA, wherein one of the

strands of the double-stranded siNA molecule is an antisense strand which comprises nucleotide sequence that is complementary to nucleotide sequence of target RNA or a portion thereof, the other strand is a sense strand which comprises nucleotide sequence that is complementary to a nucleotide sequence of the antisense strand and wherein a majority of the pyrimidine nucleotides present in the double-stranded siNA molecule comprises a sugar modification (e.g., 2'-deoxy-2'-fluoro, 2'-O-methyl, or 2'-deoxy modifications).

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that inhibits, down-regulates, or reduces expression of a target gene or that directs cleavage of a target RNA, wherein one of the strands of the double-stranded siNA molecule is an antisense strand which comprises nucleotide sequence that is complementary to nucleotide sequence of target RNA or a portion thereof, wherein the other strand is a sense strand which comprises nucleotide sequence that is complementary to a nucleotide sequence of the antisense strand and wherein a majority of the pyrimidine nucleotides present in the double-stranded siNA molecule comprises a sugar modification (e.g., 2'-deoxy-2'-fluoro, 2'-O-methyl, or 2'-deoxy modifications).

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that inhibits, down-regulates, or reduces expression of a gene or that directs cleavage of a target RNA, wherein one of the strands of the double-stranded siNA molecule is an antisense strand which comprises nucleotide sequence that is complementary to nucleotide sequence of target RNA that encodes a protein or portion thereof, the other strand is a sense strand which comprises nucleotide sequence that is complementary to a nucleotide sequence of the antisense strand and wherein a majority of the pyrimidine nucleotides present in the double-stranded siNA molecule comprises a sugar modification. In one embodiment, each strand of the siNA molecule comprises about 15 to about 30 or more (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30 or more) nucleotides, wherein each strand comprises at least about 15 nucleotides that are complementary to the nucleotides of the other strand. In one embodiment, the siNA molecule is assembled from two oligonucleotide fragments, wherein one fragment comprises the nucleotide sequence of the antisense strand of the

siNA molecule and a second fragment comprises nucleotide sequence of the sense region of the siNA molecule. In one embodiment, the sense strand is connected to the antisense strand via a linker molecule, such as a polynucleotide linker or a non-nucleotide linker. In a further embodiment, the pyrimidine nucleotides present in the sense strand are 2'-deoxy-2'-fluoro pyrimidine nucleotides and the purine nucleotides present in the sense region are 2'-deoxy purine nucleotides. In another embodiment, the pyrimidine nucleotides present in the sense strand are 2'-deoxy-2'-fluoro pyrimidine nucleotides and the purine nucleotides present in the sense region are 2'-O-methyl purine nucleotides. In still another embodiment, the pyrimidine nucleotides present in the antisense strand are 2'-deoxy-2'-fluoro pyrimidine nucleotides and any purine nucleotides present in the antisense strand are 2'-deoxy purine nucleotides. In another embodiment, the antisense strand comprises one or more 2'-deoxy-2'-fluoro pyrimidine nucleotides and one or more 2'-O-methyl purine nucleotides. In another embodiment, the pyrimidine nucleotides present in the antisense strand are 2'-deoxy-2'-fluoro pyrimidine nucleotides and any purine nucleotides present in the antisense strand are 2'-O-methyl purine nucleotides. In a further embodiment the sense strand comprises a 3'-end and a 5'-end, wherein a terminal cap moiety (e.g., an inverted deoxy abasic moiety or inverted deoxy nucleotide moiety such as inverted thymidine) is present at the 5'-end, the 3'-end, or both of the 5' and 3' ends of the sense strand. In another embodiment, the antisense strand comprises a phosphorothioate internucleotide linkage at the 3' end of the antisense strand. In another embodiment, the antisense strand comprises a glyceryl modification at the 3' end. In another embodiment, the 5'-end of the antisense strand optionally includes a phosphate group.

In any of the above-described embodiments of a double-stranded short interfering nucleic acid (siNA) molecule that inhibits expression of a target gene or that directs cleavage of a target RNA, wherein a majority of the pyrimidine nucleotides present in the double-stranded siNA molecule comprises a sugar modification, each of the two strands of the siNA molecule can comprise about 15 to about 30 or more (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30 or more) nucleotides. In one embodiment, about 15 to about 30 or more (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30 or more) nucleotides of each strand of the siNA molecule are base-paired to the complementary nucleotides of the other strand of the siNA molecule.

In another embodiment, about 15 to about 30 or more (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30 or more) nucleotides of each strand of the siNA molecule are base-paired to the complementary nucleotides of the other strand of the siNA molecule, wherein at least two 3' terminal nucleotides of each strand of the siNA molecule are not base-paired to the nucleotides of the other strand of the siNA molecule. In another embodiment, each of the two 3' terminal nucleotides of each fragment of the siNA molecule is a 2'-deoxy-pyrimidine, such as 2'-deoxy-thymidine. In one embodiment, each strand of the siNA molecule is base-paired to the complementary nucleotides of the other strand of the siNA molecule. In one embodiment, about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides of the antisense strand are base-paired to the nucleotide sequence of the target RNA or a portion thereof. In one embodiment, about 18 to about 25 (e.g., about 18, 19, 20, 21, 22, 23, 24, or 25) nucleotides of the antisense strand are base-paired to the nucleotide sequence of the target RNA or a portion thereof.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that inhibits expression of a target gene or that directs cleavage of a target RNA, wherein one of the strands of the double-stranded siNA molecule is an antisense strand which comprises nucleotide sequence that is complementary to nucleotide sequence of target RNA or a portion thereof, the other strand is a sense strand which comprises nucleotide sequence that is complementary to a nucleotide sequence of the antisense strand and wherein a majority of the pyrimidine nucleotides present in the double-stranded siNA molecule comprises a sugar modification, and wherein the 5'-end of the antisense strand optionally includes a phosphate group.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that inhibits expression of a target gene or that directs cleavage of a target RNA, wherein one of the strands of the double-stranded siNA molecule is an antisense strand which comprises nucleotide sequence that is complementary to nucleotide sequence of target RNA or a portion thereof, the other strand is a sense strand which comprises nucleotide sequence that is complementary to a nucleotide sequence of the antisense strand and wherein a majority of the pyrimidine

nucleotides present in the double-stranded siNA molecule comprises a sugar modification, and wherein the nucleotide sequence or a portion thereof of the antisense strand is complementary to a nucleotide sequence of the untranslated region or a portion thereof of the target RNA.

- 5 In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that inhibits expression of a target gene or that directs cleavage of a target RNA, wherein one of the strands of the double-stranded siNA molecule is an antisense strand which comprises nucleotide sequence that is complementary to nucleotide sequence of target RNA or a portion thereof, wherein
10 other strand is a sense strand which comprises nucleotide sequence that is complementary to a nucleotide sequence of the antisense strand, wherein a majority of the pyrimidine nucleotides present in the double-stranded siNA molecule comprises a sugar modification, and wherein the nucleotide sequence of the antisense strand is complementary to a nucleotide sequence of the target RNA or a portion thereof that is
15 present in the target RNA.

In one embodiment, the invention features a composition comprising a siNA molecule of the invention in a pharmaceutically acceptable carrier or diluent.

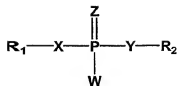
- In a non-limiting example, the introduction of chemically-modified nucleotides into nucleic acid molecules provides a powerful tool in overcoming potential limitations
20 of *in vivo* stability and bioavailability inherent to native RNA molecules that are delivered exogenously. For example, the use of chemically-modified nucleic acid molecules can enable a lower dose of a particular nucleic acid molecule for a given therapeutic effect since chemically-modified nucleic acid molecules tend to have a longer half-life in serum. Furthermore, certain chemical modifications can improve the
25 bioavailability of nucleic acid molecules by targeting particular cells or tissues and/or improving cellular up take of the nucleic acid molecule. Therefore, even if the activity of a chemically-modified nucleic acid molecule is reduced as compared to a native nucleic acid molecule, for example, when compared to an all-RNA nucleic acid molecule, the overall activity of the modified nucleic acid molecule can be greater than that of the
30 native molecule due to improved stability and/or delivery of the molecule. Unlike native

unmodified siNA, chemically-modified siNA can also minimize the possibility of activating interferon activity in humans.

In any of the embodiments of siNA molecules described herein, the antisense region of a siNA molecule of the invention can comprise a phosphorothioate internucleotide linkage at the 3'-end of said antisense region. In any of the embodiments of siNA molecules described herein, the antisense region can comprise about one to about five phosphorothioate internucleotide linkages at the 5'-end of said antisense region. In any of the embodiments of siNA molecules described herein, the 3'-terminal nucleotide overhangs of a siNA molecule of the invention can comprise ribonucleotides or deoxyribonucleotides that are chemically-modified at a nucleic acid sugar, base, or backbone. In any of the embodiments of siNA molecules described herein, the 3'-terminal nucleotide overhangs can comprise one or more universal base ribonucleotides. In any of the embodiments of siNA molecules described herein, the 3'-terminal nucleotide overhangs can comprise one or more acyclic nucleotides.

One embodiment of the invention provides an expression vector comprising a nucleic acid sequence encoding at least one siNA molecule of the invention in a manner that allows expression of the nucleic acid molecule. Another embodiment of the invention provides a mammalian cell comprising such an expression vector. The mammalian cell can be a human cell. The siNA molecule of the expression vector can comprise a sense region and an antisense region. The antisense region can comprise sequence complementary to a RNA or DNA sequence encoding the target and the sense region can comprise sequence complementary to the antisense region. The siNA molecule can comprise two distinct strands having complementary sense and antisense regions. The siNA molecule can comprise a single strand having complementary sense and antisense regions.

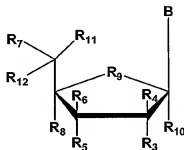
In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) against a target polynucleotide (e.g., DNA or RNA) inside a cell or reconstituted *in vitro* system, wherein the chemical modification comprises one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) nucleotides comprising a backbone modified internucleotide linkage having Formula I:



wherein each R1 and R2 is independently any nucleotide, non-nucleotide, or polynucleotide which can be naturally-occurring or chemically-modified, each X and Y is independently O, S, N, alkyl, or substituted alkyl, each Z and W is independently O, S, N, alkyl, substituted alkyl, O-alkyl, S-alkyl, alkaryl, aralkyl, or acetyl and wherein W, X, Y, and Z are optionally not all O. In another embodiment, a backbone modification of the invention comprises a phosphonoacetate and/or thiophosphonoacetate internucleotide linkage (see for example Sheehan et al., 2003, Nucleic Acids Research, 31, 4109-4118).

The chemically-modified internucleotide linkages having Formula I, for example, wherein any Z, W, X, and/or Y independently comprises a sulphur atom, can be present in one or both oligonucleotide strands of the siNA duplex, for example, in the sense strand, the antisense strand, or both strands. The siNA molecules of the invention can comprise one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) chemically-modified internucleotide linkages having Formula I at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the sense strand, the antisense strand, or both strands. For example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) chemically-modified internucleotide linkages having Formula I at the 5'-end of the sense strand, the antisense strand, or both strands. In another non-limiting example, an exemplary siNA molecule of the invention can comprise one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) pyrimidine nucleotides with chemically-modified internucleotide linkages having Formula I in the sense strand, the antisense strand, or both strands. In yet another non-limiting example, an exemplary siNA molecule of the invention can comprise one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) purine nucleotides with chemically-modified internucleotide linkages having Formula I in the sense strand, the antisense strand, or both strands. In another embodiment, a siNA molecule of the invention having internucleotide linkage(s) of Formula I also comprises a chemically-modified nucleotide or non-nucleotide having any of Formulae I-VII.

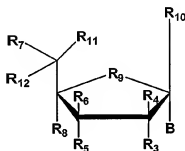
- In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) against a target polynucleotide (e.g., DNA or RNA) inside a cell or reconstituted *in vitro* system, wherein the chemical modification comprises one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) nucleotides or non-nucleotides having Formula II:



- wherein each R3, R4, R5, R6, R7, R8, R10, R11 and R12 is independently H, OH, alkyl, substituted alkyl, alkaryl or aralkyl, F, Cl, Br, CN, CF₃, OCF₃, OCN, O-alkyl, S-alkyl, N-alkyl, O-alkenyl, S-alkenyl, N-alkenyl, SO-alkyl, alkyl-OSH, alkyl-OH, O-alkyl-OH, O-alkyl-SH, S-alkyl-OH, S-alkyl-SH, alkyl-S-alkyl, alkyl-O-alkyl, ONO₂, NO₂, N₃, NH₂, aminoalkyl, aminoacid, aminoacyl, ONH₂, O-aminoalkyl, O-aminoacid, O-aminoacyl, heterocycloalkyl, heterocycloalkaryl, aminoalkylamino, polyalkylamino, substituted silyl, or group having Formula I or II; R9 is O, S, CH₂, S=O, CHF, or CF₂, and B is a nucleosidic base such as adenine, guanine, uracil, cytosine, thymine, 2-aminoadenosine, 5-methylcytosine, 2,6-diaminopurine, or any other non-naturally occurring base that can be complementary or non-complementary to target RNA or a non-nucleosidic base such as phenyl, naphthyl, 3-nitropyrrole, 5-nitroindole, nebularine, pyridone, pyridinone, or any other non-naturally occurring universal base that can be complementary or non-complementary to target RNA.
- The chemically-modified nucleotide or non-nucleotide of Formula II can be present in one or both oligonucleotide strands of the siNA duplex, for example in the sense strand, the antisense strand, or both strands. The siNA molecules of the invention can comprise one or more chemically-modified nucleotides or non-nucleotides of Formula II at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the sense strand, the antisense strand, or both strands. For example, an exemplary siNA molecule of the

invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) chemically-modified nucleotides or non-nucleotides of Formula II at the 5'-end of the sense strand, the antisense strand, or both strands. In another non-limiting example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) chemically-modified nucleotides or non-nucleotides of Formula II at the 3'-end of the sense strand, the antisense strand, or both strands.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) against a target polynucleotide (e.g., DNA or RNA) inside a cell or reconstituted *in vitro* system, wherein the chemical modification comprises one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) nucleotides or non-nucleotides having Formula III:

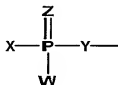


wherein each R3, R4, R5, R6, R7, R8, R10, R11 and R12 is independently H, OH, alkyl, substituted alkyl, alkaryl or aralkyl, F, Cl, Br, CN, CF3, OCF3, OCN, O-alkyl, S-alkyl, N-alkyl, O-alkenyl, S-alkenyl, N-alkenyl, SO-alkyl, alkyl-OSH, alkyl-OH, O-alkyl-OH, O-alkyl-SH, S-alkyl-OH, S-alkyl-SH, alkyl-S-alkyl, alkyl-O-alkyl, ONO2, NO2, N3, NH2, aminoalkyl, aminoacid, aminoacyl, ONH2, O-aminoalkyl, O-aminoacid, O-aminoacyl, heterocycloalkyl, heterocycloalkaryl, aminoalkylamino, polyalkylamino, substituted silyl, or group having Formula I or II; R9 is O, S, CH2, S=O, CHF, or CF2, and B is a nucleosidic base such as adenine, guanine, uracil, cytosine, thymine, 2-aminoadenosine, 5-methylcytosine, 2,6-diaminopurine, or any other non-naturally occurring base that can be employed to be complementary or non-complementary to target RNA or a non-nucleosidic base such as phenyl, naphthyl, 3-nitropyrrole, 5-nitroindole, nebularine, pyridone, pyridinone, or any other non-naturally occurring universal base that can be complementary or non-complementary to target RNA.

The chemically-modified nucleotide or non-nucleotide of Formula III can be present in one or both oligonucleotide strands of the siNA duplex, for example, in the sense strand, the antisense strand, or both strands. The siNA molecules of the invention can comprise one or more chemically-modified nucleotides or non-nucleotides of Formula III at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the sense strand, the antisense strand, or both strands. For example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) chemically-modified nucleotide(s) or non-nucleotide(s) of Formula III at the 5'-end of the sense strand, the antisense strand, or both strands. In another non-limiting example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) chemically-modified nucleotide or non-nucleotide of Formula III at the 3'-end of the sense strand, the antisense strand, or both strands.

In another embodiment, a siNA molecule of the invention comprises a nucleotide having Formula II or III, wherein the nucleotide having Formula II or III is in an inverted configuration. For example, the nucleotide having Formula II or III is connected to the siNA construct in a 3'-3', 3'-2', 2'-3', or 5'-5' configuration, such as at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of one or both siNA strands.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) against a target polynucleotide (e.g., DNA or RNA) inside a cell or reconstituted *in vitro* system, wherein the chemical modification comprises a 5'-terminal phosphate group having Formula IV:



wherein each X and Y is independently O, S, N, alkyl, substituted alkyl, or alkylhalo; wherein each Z and W is independently O, S, N, alkyl, substituted alkyl, O-alkyl, S-alkyl, alkaryl, aralkyl, alkylhalo, or acetyl; and wherein W, X, Y and Z are not all O.

In one embodiment, the invention features a siNA molecule having a 5'-terminal phosphate group having Formula IV on the target-complementary strand, for example, a strand complementary to a target RNA, wherein the siNA molecule comprises an all RNA siNA molecule. In another embodiment, the invention features a siNA molecule
5 having a 5'-terminal phosphate group having Formula IV on the target-complementary strand wherein the siNA molecule also comprises about 1 to about 3 (e.g., about 1, 2, or 3) nucleotide 3'-terminal nucleotide overhangs having about 1 to about 4 (e.g., about 1, 2, 3, or 4) deoxyribonucleotides on the 3'-end of one or both strands. In another embodiment, a 5'-terminal phosphate group having Formula IV is present on the target-
10 complementary strand of a siNA molecule of the invention, for example a siNA molecule having chemical modifications having any of Formulae I-VII.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) against a target polynucleotide (e.g., DNA or RNA) inside a cell or reconstituted *in vitro* system,
15 wherein the chemical modification comprises one or more phosphorothioate internucleotide linkages. For example, in a non-limiting example, the invention features a chemically-modified short interfering nucleic acid (siNA) having about 1, 2, 3, 4, 5, 6, 7, 8 or more phosphorothioate internucleotide linkages in one siNA strand. In yet another embodiment, the invention features a chemically-modified short interfering
20 nucleic acid (siNA) individually having about 1, 2, 3, 4, 5, 6, 7, 8 or more phosphorothioate internucleotide linkages in both siNA strands. The phosphorothioate internucleotide linkages can be present in one or both oligonucleotide strands of the siNA duplex, for example in the sense strand, the antisense strand, or both strands. The siNA molecules of the invention can comprise one or more phosphorothioate
25 internucleotide linkages at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the sense strand, the antisense strand, or both strands. For example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) consecutive phosphorothioate internucleotide linkages at the 5'-end of the sense strand, the antisense strand, or both strands. In another non-limiting example, an
30 exemplary siNA molecule of the invention can comprise one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) pyrimidine phosphorothioate internucleotide linkages in the sense strand, the antisense strand, or both strands. In yet another non-limiting example,

an exemplary siRNA molecule of the invention can comprise one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) purine phosphorothioate internucleotide linkages in the sense strand, the antisense strand, or both strands.

In one embodiment, a siRNA molecule of the invention is featured, wherein the
5 sense strand comprises one or more, for example, about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or about one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of
10 the 3'- and 5'-ends of the sense strand; and wherein the antisense strand comprises about 1 to about 10 or more, specifically about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and
15 optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the antisense strand. In another embodiment, one or more, for example about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more, pyrimidine nucleotides of the sense and/or antisense siRNA strand are chemically-modified with 2'-deoxy, 2'-O-methyl and/or 2'-deoxy-2'-fluoro nucleotides, with or without one or more, for example about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or
20 more, phosphorothioate internucleotide linkages and/or a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends, being present in the same or different strand.

In another embodiment, a siRNA molecule of the invention is featured, wherein the sense strand comprises about 1 to about 5, specifically about 1, 2, 3, 4, or 5
25 phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the sense strand; and wherein the antisense strand comprises about 1 to about 5 or more, specifically about 1,
30 2, 3, 4, 5, or more phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro,

and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the antisense strand. In another embodiment, one or more, for example about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more, pyrimidine nucleotides of the sense and/or antisense siNA strand are chemically-modified with 2'-deoxy, 2'-O-methyl and/or 2'-deoxy-2'-fluoro nucleotides, with or without about 1 to about 5 or more, for example about 1, 2, 3, 4, 5, or more phosphorothioate internucleotide linkages and/or a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends, being present in the same or different strand.

10 In one embodiment, a siNA molecule of the invention is featured, wherein the antisense strand comprises one or more, for example, about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more phosphorothioate internucleotide linkages, and/or about one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the sense strand; and wherein the antisense strand comprises about 1 to about 10 or more, specifically about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the antisense strand. In another embodiment, one or more, for example about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more pyrimidine nucleotides of the sense and/or antisense siNA strand are chemically-modified with 2'-deoxy, 2'-O-methyl and/or 2'-deoxy-2'-fluoro nucleotides, with or without one or more, for example, about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more phosphorothioate internucleotide linkages and/or a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3' and 5'-ends, being present in the same or different strand.

In another embodiment, a siNA molecule of the invention is featured, wherein the antisense strand comprises about 1 to about 5 or more, specifically about 1, 2, 3, 4, 5 or more phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more

(e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the sense strand; and wherein the antisense strand comprises about 1 to about 5 or more, specifically about 1, 2, 3, 4, 5 or more phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the antisense strand. In another embodiment, one or more, for example about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more pyrimidine nucleotides of the sense and/or antisense siNA strand are chemically-modified with 2'-deoxy, 2'-O-methyl and/or 2'-deoxy-2'-fluoro nucleotides, with or without about 1 to about 5, for example about 1, 2, 3, 4, 5 or more phosphorothioate internucleotide linkages and/or a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends, being present in the same or different strand.

In one embodiment, a chemically-modified short interfering nucleic acid (siNA) molecule of the invention comprises about 1 to about 5 or more (specifically about 1, 2, 3, 4, 5 or more) phosphorothioate internucleotide linkages in each strand of the siNA molecule.

In another embodiment, a siNA molecule of the invention comprises 2'-5' internucleotide linkages. The 2'-5' internucleotide linkage(s) can be at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of one or both siNA sequence strands. In addition, the 2'-5' internucleotide linkage(s) can be present at various other positions within one or both siNA sequence strands, for example, about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more including every internucleotide linkage of a pyrimidine nucleotide in one or both strands of the siNA molecule can comprise a 2'-5' internucleotide linkage, or about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more including every internucleotide linkage of a purine nucleotide in one or both strands of the siNA molecule can comprise a 2'-5' internucleotide linkage.

In another embodiment, a chemically-modified siNA molecule of the invention comprises a duplex having two strands, one or both of which can be chemically-modified, wherein each strand is independently about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides in length, wherein the

duplex has about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) base pairs, and wherein the chemical modification comprises a structure having any of Formulae I-VII. For example, an exemplary chemically-modified siNA molecule of the invention comprises a duplex having two strands, one or both of which can be chemically-modified with a chemical modification having any of Formulae I-VII or any combination thereof, wherein each strand consists of about 21 nucleotides, each having a 2-nucleotide 3'-terminal nucleotide overhang, and wherein the duplex has about 19 base pairs. In another embodiment, a siNA molecule of the invention comprises a single stranded hairpin structure, wherein the siNA is about 36 to about 70 (e.g., about 36, 40, 45, 50, 55, 60, 65, or 70) nucleotides in length having about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) base pairs, and wherein the siNA can include a chemical modification comprising a structure having any of Formulae I-VII or any combination thereof. For example, an exemplary chemically-modified siNA molecule of the invention comprises a linear oligonucleotide having about 42 to about 50 (e.g., about 42, 43, 44, 45, 46, 47, 48, 49, or 50) nucleotides that is chemically-modified with a chemical modification having any of Formulae I-VII or any combination thereof, wherein the linear oligonucleotide forms a hairpin structure having about 19 to about 21 (e.g., 19, 20, or 21) base pairs and a 2-nucleotide 3'-terminal nucleotide overhang. In another embodiment, a linear hairpin siNA molecule of the invention contains a stem loop motif, wherein the loop portion of the siNA molecule is biodegradable. For example, a linear hairpin siNA molecule of the invention is designed such that degradation of the loop portion of the siNA molecule *in vivo* can generate a double-stranded siNA molecule with 3'-terminal overhangs, such as 3'-terminal nucleotide overhangs comprising about 2 nucleotides.

In another embodiment, a siNA molecule of the invention comprises a hairpin structure, wherein the siNA is about 25 to about 50 (e.g., about 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, or 50) nucleotides in length having about 3 to about 25 (e.g., about 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, or 25) base pairs, and wherein the siNA can include one or more chemical modifications comprising a structure having any of Formulae I-VII or any combination thereof. For example, an exemplary chemically-modified siNA molecule of the invention comprises a linear oligonucleotide having about 25 to about 35 (e.g., about

25, 26, 27, 28, 29, 30, 31, 32, 33, 34, or 35) nucleotides that is chemically-modified with one or more chemical modifications having any of Formulae I-VII or any combination thereof, wherein the linear oligonucleotide forms a hairpin structure having about 3 to about 25 (e.g., about 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, or 25) base pairs and a 5'-terminal phosphate group that can be chemically modified as described herein (for example a 5'-terminal phosphate group having Formula IV). In another embodiment, a linear hairpin siNA molecule of the invention contains a stem loop motif, wherein the loop portion of the siNA molecule is biodegradable. In one embodiment, a linear hairpin siNA molecule of the invention comprises a loop portion comprising a non-nucleotide linker.

In another embodiment, a siNA molecule of the invention comprises an asymmetric hairpin structure, wherein the siNA is about 25 to about 50 (e.g., about 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, or 50) nucleotides in length having about 3 to about 25 (e.g., about 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, or 25) base pairs, and wherein the siNA can include one or more chemical modifications comprising a structure having any of Formulae I-VII or any combination thereof. For example, an exemplary chemically-modified siNA molecule of the invention comprises a linear oligonucleotide having about 25 to about 35 (e.g., about 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, or 35) nucleotides that is chemically-modified with one or more chemical modifications having any of Formulae I-VII or any combination thereof, wherein the linear oligonucleotide forms an asymmetric hairpin structure having about 3 to about 25 (e.g., about 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, or 25) base pairs and a 5'-terminal phosphate group that can be chemically modified as described herein (for example a 5'-terminal phosphate group having Formula IV). In one embodiment, an asymmetric hairpin siNA molecule of the invention contains a stem loop motif, wherein the loop portion of the siNA molecule is biodegradable. In another embodiment, an asymmetric hairpin siNA molecule of the invention comprises a loop portion comprising a non-nucleotide linker.

In another embodiment, a siNA molecule of the invention comprises an asymmetric double stranded structure having separate polynucleotide strands comprising

sense and antisense regions, wherein the antisense region is about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides in length, wherein the sense region is about 3 to about 25 (e.g., about 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, or 25) nucleotides in length, wherein

5 the sense region and the antisense region have at least 3 complementary nucleotides, and wherein the siNA can include one or more chemical modifications comprising a structure having any of Formulae I-VII or any combination thereof. For example, an exemplary chemically-modified siNA molecule of the invention comprises an asymmetric double stranded structure having separate polynucleotide strands comprising sense and antisense

10 regions, wherein the antisense region is about 18 to about 23 (e.g., about 18, 19, 20, 21, 22, or 23) nucleotides in length and wherein the sense region is about 3 to about 15 (e.g., about 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, or 15) nucleotides in length, wherein the sense region the antisense region have at least 3 complementary nucleotides, and wherein the siNA can include one or more chemical modifications comprising a structure having any

15 of Formulae I-VII or any combination thereof. In another embodiment, the asymmetric double stranded siNA molecule can also have a 5'-terminal phosphate group that can be chemically modified as described herein (for example a 5'-terminal phosphate group having Formula IV).

In another embodiment, a siNA molecule of the invention comprises a circular

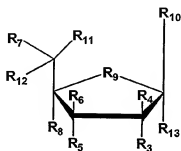
20 nucleic acid molecule, wherein the siNA is about 38 to about 70 (e.g., about 38, 40, 45, 50, 55, 60, 65, or 70) nucleotides in length having about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) base pairs, and wherein the siNA can include a chemical modification, which comprises a structure having any of Formulae I-VII or any combination thereof. For example, an exemplary chemically-

25 modified siNA molecule of the invention comprises a circular oligonucleotide having about 42 to about 50 (e.g., about 42, 43, 44, 45, 46, 47, 48, 49, or 50) nucleotides that is chemically-modified with a chemical modification having any of Formulae I-VII or any combination thereof, wherein the circular oligonucleotide forms a dumbbell shaped structure having about 19 base pairs and 2 loops.

30 In another embodiment, a circular siNA molecule of the invention contains two loop motifs, wherein one or both loop portions of the siNA molecule is biodegradable.

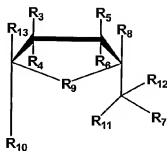
For example, a circular siNA molecule of the invention is designed such that degradation of the loop portions of the siNA molecule *in vivo* can generate a double-stranded siNA molecule with 3'-terminal overhangs, such as 3'-terminal nucleotide overhangs comprising about 2 nucleotides.

- 5 In one embodiment, a siNA molecule of the invention comprises at least one (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) abasic moiety, for example a compound having Formula V:



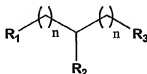
- wherein each R3, R4, R5, R6, R7, R8, R10, R11, R12, and R13 is independently H, OH,
 10 alkyl, substituted alkyl, alkaryl or aralkyl, F, Cl, Br, CN, CF3, OCF3, OCN, O-alkyl, S-alkyl, N-alkyl, O-alkenyl, S-alkenyl, N-alkenyl, SO-alkyl, alkyl-OSH, alkyl-OH, O-alkyl-OH, O-alkyl-SH, S-alkyl-OH, S-alkyl-SH, alkyl-S-alkyl, alkyl-O-alkyl, ONO2, NO2, N3, NH2, aminoalkyl, aminoacid, aminoacyl, ONH2, O-aminoalkyl, O-aminoacid, O-aminoacyl, heterocycloalkyl, heterocycloalkaryl, aminoalkylamino, polyalkylamino,
 15 substituted silyl, or group having Formula I or II; R9 is O, S, CH2, S=O, CHF, or CF2.

In one embodiment, a siNA molecule of the invention comprises at least one (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) inverted abasic moiety, for example a compound having Formula VI:



wherein each R3, R4, R5, R6, R7, R8, R10, R11, R12, and R13 is independently H, OH, alkyl, substituted alkyl, alkaryl or aralkyl, F, Cl, Br, CN, CF3, OCF3, OCN, O-alkyl, S-alkyl, N-alkyl, O-alkenyl, S-alkenyl, N-alkenyl, SO-alkyl, alkyl-OSH, alkyl-OH, O-alkyl-OH, O-alkyl-SH, S-alkyl-OH, S-alkyl-SH, alkyl-S-alkyl, alkyl-O-alkyl, ONO2, NO2, N3, NH2, aminoalkyl, aminoacid, aminoacyl, ONH2, O-aminoalkyl, O-aminoacid, O-aminoacyl, heterocycloalkyl, heterocycloalkaryl, aminoalkylamino, polyalkylamino, substituted silyl, or group having Formula I or II; R9 is O, S, CH2, S=O, CHF, or CF2, and either R2, R3, R8 or R13 serve as points of attachment to the siNA molecule of the invention.

10 In another embodiment, a siNA molecule of the invention comprises at least one (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) substituted polyalkyl moieties, for example a compound having Formula VII:



wherein each n is independently an integer from 1 to 12, each R1, R2 and R3 is independently H, OH, alkyl, substituted alkyl, alkaryl or aralkyl, F, Cl, Br, CN, CF3, OCF3, OCN, O-alkyl, S-alkyl, N-alkyl, O-alkenyl, S-alkenyl, N-alkenyl, SO-alkyl, alkyl-OSH, alkyl-OH, O-alkyl-OH, O-alkyl-SH, S-alkyl-OH, S-alkyl-SH, alkyl-S-alkyl, alkyl-O-alkyl, ONO2, NO2, N3, NH2, aminoalkyl, aminoacid, aminoacyl, ONH2, O-aminoalkyl, O-aminoacid, O-aminoacyl, heterocycloalkyl, heterocycloalkaryl, aminoalkylamino, polyalkylamino, substituted silyl, or a group having Formula I, and R1, R2 or R3 serves as points of attachment to the siNA molecule of the invention.

In another embodiment, a siNA molecule of the invention comprises a compound having Formula VII, wherein R1 and R2 are hydroxyl (OH) groups, n = 1, and R3 comprises O and is the point of attachment to the 3'-end, the 5'-end, or both of the 3' and 5'-ends of one or both strands of a double-stranded siNA molecule of the invention or to a single-stranded siNA molecule of the invention. This modification is referred to herein as "glyceryl" (for example modification 6 in Figure 10).

In another embodiment, a chemically modified nucleoside or non-nucleoside (e.g. a moiety having any of Formula V, VI or VII) of the invention is at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of a siNA molecule of the invention. For example, chemically modified nucleoside or non-nucleoside (e.g., a moiety having Formula V, VI or VII) can be present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense strand, the sense strand, or both antisense and sense strands of the siNA molecule. In one embodiment, the chemically modified nucleoside or non-nucleoside (e.g., a moiety having Formula V, VI or VII) is present at the 5'-end and 3'-end of the sense strand and the 3'-end of the antisense strand of a double stranded siNA molecule of the invention. In one embodiment, the chemically modified nucleoside or non-nucleoside (e.g., a moiety having Formula V, VI or VII) is present at the terminal position of the 5'-end and 3'-end of the sense strand and the 3'-end of the antisense strand of a double stranded siNA molecule of the invention. In one embodiment, the chemically modified nucleoside or non-nucleoside (e.g., a moiety having Formula V, VI or VII) is present at the two terminal positions of the 5'-end and 3'-end of the sense strand and the 3'-end of the antisense strand of a double stranded siNA molecule of the invention. In one embodiment, the chemically modified nucleoside or non-nucleoside (e.g., a moiety having Formula V, VI or VII) is present at the penultimate position of the 5'-end and 3'-end of the sense strand and the 3'-end of the antisense strand of a double stranded siNA molecule of the invention. In addition, a moiety having Formula VII can be present at the 3'-end or the 5'-end of a hairpin siNA molecule as described herein.

In another embodiment, a siNA molecule of the invention comprises an abasic residue having Formula V or VI, wherein the abasic residue having Formula VI or VI is connected to the siNA construct in a 3'-3', 3'-2', 2'-3', or 5'-5' configuration, such as at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of one or both siNA strands.

In one embodiment, a siNA molecule of the invention comprises one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) locked nucleic acid (LNA) nucleotides, for example, at the 5'-end, the 3'-end, both of the 5' and 3'-ends, or any combination thereof, of the siNA molecule.

In another embodiment, a siNA molecule of the invention comprises one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) acyclic nucleotides, for example, at the

5'-end, the 3'-end, both of the 5' and 3'-ends, or any combination thereof, of the siNA molecule.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention comprising a sense region, wherein any
 5 (e.g., one or more or all) pyrimidine nucleotides present in the sense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any (e.g., one or more or all)
 10 purine nucleotides present in the sense region are 2'-deoxy purine nucleotides (e.g., wherein all purine nucleotides are 2'-deoxy purine nucleotides or alternately a plurality of purine nucleotides are 2'-deoxy purine nucleotides).

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention comprising a sense region, wherein any
 (e.g., one or more or all) pyrimidine nucleotides present in the sense region are 2'-deoxy-
 15 2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any (e.g., one or more or all)
 purine nucleotides present in the sense region are 2'-deoxy purine nucleotides (e.g., wherein all purine nucleotides are 2'-deoxy purine nucleotides or alternately a plurality
 20 of purine nucleotides are 2'-deoxy purine nucleotides), wherein any nucleotides comprising a 3'-terminal nucleotide overhang that are present in said sense region are 2'-deoxy nucleotides.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention comprising a sense region, wherein any
 25 (e.g., one or more or all) pyrimidine nucleotides present in the sense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any (e.g., one or more or all)
 purine nucleotides present in the sense region are 2'-O-methyl purine nucleotides (e.g.,
 30 wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides).

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention comprising a sense region, wherein any (e.g., one or more or all) pyrimidine nucleotides present in the sense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), wherein any (e.g., one or more or all) purine nucleotides present in the sense region are 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides), and wherein any nucleotides comprising a 3'-terminal nucleotide overhang that are present in said sense region are 2'-deoxy nucleotides.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention comprising an antisense region, wherein any (e.g., one or more or all) pyrimidine nucleotides present in the antisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any (e.g., one or more or all) purine nucleotides present in the antisense region are 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides).

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention comprising an antisense region, wherein any (e.g., one or more or all) pyrimidine nucleotides present in the antisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), wherein any (e.g., one or more or all) purine nucleotides present in the antisense region are 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides), and wherein any nucleotides comprising a 3'-terminal nucleotide overhang that are present in said antisense region are 2'-deoxy nucleotides.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention comprising an antisense region, wherein any (*e.g.*, one or more or all) pyrimidine nucleotides present in the antisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (*e.g.*, wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any (*e.g.*, one or more or all) purine nucleotides present in the antisense region are 2'-deoxy purine nucleotides (*e.g.*, wherein all purine nucleotides are 2'-deoxy purine nucleotides or alternately a plurality of purine nucleotides are 2'-deoxy purine nucleotides).

10 In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention comprising an antisense region, wherein any (*e.g.*, one or more or all) pyrimidine nucleotides present in the antisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (*e.g.*, wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any (*e.g.*, one or more or all) purine nucleotides present in the antisense region are 2'-O-methyl purine nucleotides (*e.g.*, wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides).

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention capable of mediating RNA interference (RNAi) against a target polynucleotide (*e.g.*, DNA or RNA) inside a cell or reconstituted *in vitro* system comprising a sense region, wherein one or more pyrimidine nucleotides present in the sense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (*e.g.*, wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and one or more purine nucleotides present in the sense region are 2'-deoxy purine nucleotides (*e.g.*, wherein all purine nucleotides are 2'-deoxy purine nucleotides or alternately a plurality of purine nucleotides are 2'-deoxy purine nucleotides), and an antisense region, wherein one or more pyrimidine nucleotides present in the antisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (*e.g.*, wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of

pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and one or more purine nucleotides present in the antisense region are 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides). The sense region and/or the antisense region can have a terminal cap modification, such as any modification described herein or shown in **Figure 10**, that is optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the sense and/or antisense sequence. The sense and/or antisense region can optionally further comprise a 3'-terminal nucleotide overhang having about 1 to about 4 (e.g., about 1, 2, 3, or 4) 2'-deoxynucleotides. The overhang nucleotides can further comprise one or more (e.g., about 1, 2, 3, 4 or more) phosphorothioate, phosphonoacetate, and/or thiophosphonoacetate internucleotide linkages. Non-limiting examples of these chemically-modified siNAs are shown in **Figures 4 and 5 and Table II** herein. In any of these described embodiments, the purine nucleotides present in the sense region are alternatively 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides) and one or more purine nucleotides present in the antisense region are 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides). Also, in any of these embodiments, one or more purine nucleotides present in the sense region are alternatively purine ribonucleotides (e.g., wherein all purine nucleotides are purine ribonucleotides or alternately a plurality of purine nucleotides are purine ribonucleotides) and any purine nucleotides present in the antisense region are 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides). Additionally, in any of these embodiments, one or more purine nucleotides present in the sense region and/or present in the antisense region are alternatively selected from the group consisting of 2'-deoxy nucleotides, locked nucleic acid (LNA) nucleotides, 2'-methoxyethyl nucleotides, 4'-thionucleotides, and 2'-O-methyl nucleotides (e.g., wherein all purine nucleotides are selected from the group consisting of 2'-deoxy nucleotides, locked nucleic acid (LNA) nucleotides, 2'-methoxyethyl nucleotides, 4'-thionucleotides, and 2'-O-methyl nucleotides or alternately a plurality of purine

nucleotides are selected from the group consisting of 2'-deoxy nucleotides, locked nucleic acid (LNA) nucleotides, 2'-methoxyethyl nucleotides, 4'-thionucleotides, and 2'-O-methyl nucleotides).

In another embodiment, any modified nucleotides present in the siRNA molecules of the invention, preferably in the antisense strand of the siRNA molecules of the invention, but also optionally in the sense and/or both antisense and sense strands, comprise modified nucleotides having properties or characteristics similar to naturally occurring ribonucleotides. For example, the invention features siRNA molecules including modified nucleotides having a Northern conformation (e.g., Northern pseudorotation cycle, see for example Saenger, *Principles of Nucleic Acid Structure*, Springer-Verlag ed., 1984). As such, chemically modified nucleotides present in the siRNA molecules of the invention, preferably in the antisense strand of the siRNA molecules of the invention, but also optionally in the sense and/or both antisense and sense strands, are resistant to nuclease degradation while at the same time maintaining the capacity to mediate RNAi. Non-limiting examples of nucleotides having a northern configuration include locked nucleic acid (LNA) nucleotides (e.g., 2'-O, 4'-C-methylene-(D-ribofuranosyl) nucleotides); 2'-methoxyethoxy (MOE) nucleotides; 2'-methyl-thio-ethyl, 2'-deoxy-2'-fluoro nucleotides, 2'-deoxy-2'-chloro nucleotides, 2'-azido nucleotides, and 2'-O-methyl nucleotides.

In one embodiment, the sense strand of a double stranded siRNA molecule of the invention comprises a terminal cap moiety, (see for example **Figure 10**) such as an inverted deoxyribose moiety, at the 3'-end, 5'-end, or both 3' and 5'-ends of the sense strand.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid molecule (siRNA) capable of mediating RNA interference (RNAi) against a target polynucleotide (e.g., DNA or RNA) inside a cell or reconstituted *in vitro* system, wherein the chemical modification comprises a conjugate covalently attached to the chemically-modified siRNA molecule. Non-limiting examples of conjugates contemplated by the invention include conjugates and ligands described in Vargeese *et al.*, USSN 10/427,160, filed April 30, 2003, incorporated by reference herein in its entirety, including the drawings. In another embodiment, the conjugate is covalently

attached to the chemically-modified siNA molecule via a biodegradable linker. In one embodiment, the conjugate molecule is attached at the 3'-end of either the sense strand, the antisense strand, or both strands of the chemically-modified siNA molecule. In another embodiment, the conjugate molecule is attached at the 5'-end of either the sense strand, the antisense strand, or both strands of the chemically-modified siNA molecule. In yet another embodiment, the conjugate molecule is attached both the 3'-end and 5'-end of either the sense strand, the antisense strand, or both strands of the chemically-modified siNA molecule, or any combination thereof. In one embodiment, a conjugate molecule of the invention comprises a molecule that facilitates delivery of a chemically-modified siNA molecule into a biological system, such as a cell. In another embodiment, the conjugate molecule attached to the chemically-modified siNA molecule is a polyethylene glycol, human serum albumin, or a ligand for a cellular receptor that can mediate cellular uptake. Examples of specific conjugate molecules contemplated by the instant invention that can be attached to chemically-modified siNA molecules are described in Vargeese *et al.*, U.S. Serial No. 10/201,394, filed July 22, 2002 incorporated by reference herein. The type of conjugates used and the extent of conjugation of siNA molecules of the invention can be evaluated for improved pharmacokinetic profiles, bioavailability, and/or stability of siNA constructs while at the same time maintaining the ability of the siNA to mediate RNAi activity. As such, one skilled in the art can screen siNA constructs that are modified with various conjugates to determine whether the siNA conjugate complex possesses improved properties while maintaining the ability to mediate RNAi, for example in animal models as are generally known in the art.

In one embodiment, the invention features a short interfering nucleic acid (siNA) molecule of the invention, wherein the siNA further comprises a nucleotide, non-nucleotide, or mixed nucleotide/non-nucleotide linker that joins the sense region of the siNA to the antisense region of the siNA. In one embodiment, a nucleotide linker of the invention can be a linker of ≥ 2 nucleotides in length, for example about 3, 4, 5, 6, 7, 8, 9, or 10 nucleotides in length. In another embodiment, the nucleotide linker can be a nucleic acid aptamer. By "aptamer" or "nucleic acid aptamer" as used herein is meant a nucleic acid molecule that binds specifically to a target molecule wherein the nucleic acid molecule has sequence that comprises a sequence recognized by the target molecule in its natural setting. Alternately, an aptamer can be a nucleic acid molecule that binds to

- a target molecule where the target molecule does not naturally bind to a nucleic acid. The target molecule can be any molecule of interest. For example, the aptamer can be used to bind to a ligand-binding domain of a protein, thereby preventing interaction of the naturally occurring ligand with the protein. This is a non-limiting example and those in the art will recognize that other embodiments can be readily generated using techniques generally known in the art. (See, for example, Gold *et al.*, 1995, *Annu. Rev. Biochem.*, 64, 763; Brody and Gold, 2000, *J. Biotechnol.*, 74, 5; Sun, 2000, *Curr. Opin. Mol. Ther.*, 2, 100; Kusser, 2000, *J. Biotechnol.*, 74, 27; Hermann and Patel, 2000, *Science*, 287, 820; and Jayasena, 1999, *Clinical Chemistry*, 45, 1628.)
- 10 In yet another embodiment, a non-nucleotide linker of the invention comprises abasic nucleotide, polyether, polyamine, polyamide, peptide, carbohydrate, lipid, polyhydrocarbon, or other polymeric compounds (e.g. polyethylene glycols such as those having between 2 and 100 ethylene glycol units). Specific examples include those described by Seela and Kaiser, *Nucleic Acids Res.* 1990, 18:6353 and *Nucleic Acids Res.* 1987, 15:3113; Cload and Schepartz, *J. Am. Chem. Soc.* 1991, 113:6324; Richardson and Schepartz, *J. Am. Chem. Soc.* 1991, 113:5109; Ma *et al.*, *Nucleic Acids Res.* 1993, 21:2585 and *Biochemistry* 1993, 32:1751; Durand *et al.*, *Nucleic Acids Res.* 1990, 18:6353; McCurdy *et al.*, *Nucleosides & Nucleotides* 1991, 10:287; Jschke *et al.*, *Tetrahedron Lett.* 1993, 34:301; Ono *et al.*, *Biochemistry* 1991, 30:9914; Arnold *et al.*, 20 International Publication No. WO 89/02439; Usman *et al.*, International Publication No. WO 95/06731; Dudycz *et al.*, International Publication No. WO 95/11910 and Ferentz and Verdine, *J. Am. Chem. Soc.* 1991, 113:4000, all hereby incorporated by reference herein. A "non-nucleotide" further means any group or compound that can be incorporated into a nucleic acid chain in the place of one or more nucleotide units, including either sugar and/or phosphate substitutions, and allows the remaining bases to exhibit their enzymatic activity. The group or compound can be abasic in that it does not contain a commonly recognized nucleotide base, such as adenosine, guanine, cytosine, uracil or thymine, for example at the C1 position of the sugar.

- In one embodiment, the invention features a short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) inside a cell or reconstituted *in vitro* system, wherein one or both strands of the siNA molecule that are assembled from

two separate oligonucleotides do not comprise any ribonucleotides. For example, a siNA molecule can be assembled from a single oligonucleotide where the sense and antisense regions of the siNA comprise separate oligonucleotides that do not have any ribonucleotides (e.g., nucleotides having a 2'-OH group) present in the oligonucleotides.

5 In another example, a siNA molecule can be assembled from a single oligonucleotide where the sense and antisense regions of the siNA are linked or circularized by a nucleotide or non-nucleotide linker as described herein, wherein the oligonucleotide does not have any ribonucleotides (e.g., nucleotides having a 2'-OH group) present in the oligonucleotide. Applicant has surprisingly found that the presense of ribonucleotides

10 (e.g., nucleotides having a 2'-hydroxyl group) within the siNA molecule is not required or essential to support RNAi activity. As such, in one embodiment, all positions within the siNA can include chemically modified nucleotides and/or non-nucleotides such as nucleotides and or non-nucleotides having Formula I, II, III, IV, V, VI, or VII or any combination thereof to the extent that the ability of the siNA molecule to support RNAi

15 activity in a cell is maintained.

In one embodiment, a siNA molecule of the invention is a single stranded siNA molecule that mediates RNAi activity in a cell or reconstituted *in vitro* system comprising a single stranded polynucleotide having complementarity to a target nucleic acid sequence. In another embodiment, the single stranded siNA molecule of the

20 invention comprises a 5'-terminal phosphate group. In another embodiment, the single stranded siNA molecule of the invention comprises a 5'-terminal phosphate group and a 3'-terminal phosphate group (e.g., a 2',3'-cyclic phosphate). In another embodiment, the single stranded siNA molecule of the invention comprises about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides. In yet

25 another embodiment, the single stranded siNA molecule of the invention comprises one or more chemically modified nucleotides or non-nucleotides described herein. For example, all the positions within the siNA molecule can include chemically-modified nucleotides such as nucleotides having any of Formulae I-VII, or any combination thereof to the extent that the ability of the siNA molecule to support RNAi activity in a

30 cell is maintained.

In one embodiment, a siNA molecule of the invention is a single stranded siNA molecule that mediates RNAi activity in a cell or reconstituted *in vitro* system comprising a single stranded polynucleotide having complementarity to a target nucleic acid sequence, wherein one or more pyrimidine nucleotides present in the siNA are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any purine nucleotides present in the antisense region are 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides), and a terminal cap modification, such as any modification described herein or shown in **Figure 10**, that is optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense sequence. The siNA optionally further comprises about 1 to about 4 or more (e.g., about 1, 2, 3, 4 or more) terminal 2'-deoxynucleotides at the 3'-end of the siNA molecule, wherein the terminal nucleotides can further comprise one or more (e.g., 1, 2, 3, 4 or more) phosphorothioate, phosphonoacetate, and/or thiophosphonoacetate internucleotide linkages, and wherein the siNA optionally further comprises a terminal phosphate group, such as a 5'-terminal phosphate group. In any of these embodiments, any purine nucleotides present in the antisense region are alternatively 2'-deoxy purine nucleotides (e.g., wherein all purine nucleotides are 2'-deoxy purine nucleotides or alternately a plurality of purine nucleotides are 2'-deoxy purine nucleotides). Also, in any of these embodiments, any purine nucleotides present in the siNA (i.e., purine nucleotides present in the sense and/or antisense region) can alternatively be locked nucleic acid (LNA) nucleotides (e.g., wherein all purine nucleotides are LNA nucleotides or alternately a plurality of purine nucleotides are LNA nucleotides). Also, in any of these embodiments, any purine nucleotides present in the siNA are alternatively 2'-methoxyethyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-methoxyethyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-methoxyethyl purine nucleotides). In another embodiment, any modified nucleotides present in the single stranded siNA molecules of the invention comprise modified nucleotides having properties or characteristics similar to naturally occurring ribonucleotides. For example, the invention features siNA molecules including modified

nucleotides having a Northern conformation (e.g., Northern pseudorotation cycle, see for example Saenger, *Principles of Nucleic Acid Structure*, Springer-Verlag ed., 1984). As such, chemically modified nucleotides present in the single stranded siNA molecules of the invention are preferably resistant to nuclease degradation while at the same time

5 maintaining the capacity to mediate RNAi.

In one embodiment, a siNA molecule of the invention comprises chemically modified nucleotides or non-nucleotides (e.g., having any of Formulae I-VII, such as 2'-deoxy, 2'-deoxy-2'-fluoro, or 2'-O-methyl nucleotides) at alternating positions within one or more strands or regions of the siNA molecule. For example, such chemical

10 modifications can be introduced at every other position of a RNA based siNA molecule, starting at either the first or second nucleotide from the 3'-end or 5'-end of the siNA. In a non-limiting example, a double stranded siNA molecule of the invention in which each strand of the siNA is 21 nucleotides in length is featured wherein positions 1, 3, 5, 7, 9, 11, 13, 15, 17, 19 and 21 of each strand are chemically modified (e.g., with compounds

15 having any of Formulae I-VII, such as such as 2'-deoxy, 2'-deoxy-2'-fluoro, or 2'-O-methyl nucleotides). In another non-limiting example, a double stranded siNA molecule of the invention in which each strand of the siNA is 21 nucleotides in length is featured wherein positions 2, 4, 6, 8, 10, 12, 14, 16, 18, and 20 of each strand are chemically modified (e.g., with compounds having any of Formulae I-VII, such as such as 2'-deoxy,

20 2'-deoxy-2'-fluoro, or 2'-O-methyl nucleotides). Such siNA molecules can further comprise terminal cap moieties and/or backbone modifications as described herein.

In one embodiment, the invention features a method for modulating the expression of a target gene within a cell comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands

25 comprises a sequence complementary to RNA of the target gene; and (b) introducing the siNA molecule into a cell under conditions suitable to modulate the expression of the target gene in the cell.

In one embodiment, the invention features a method for modulating the expression of a target gene within a cell comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands

30 comprises a sequence complementary to RNA of the target gene and wherein the sense

sense sequence of the siRNA comprises a sequence identical or substantially similar to the sequence of the target RNA; and (b) introducing the siRNA molecule into a cell under conditions suitable to modulate the expression of the target gene in the cell.

5 In another embodiment, the invention features a method for modulating the expression of more than one target gene within a cell comprising: (a) synthesizing siRNA molecules of the invention, which can be chemically-modified, wherein one of the siRNA strands comprises a sequence complementary to RNA of the target genes; and (b) introducing the siRNA molecules into a cell under conditions suitable to modulate the expression of the target genes in the cell.

10 In another embodiment, the invention features a method for modulating the expression of two or more target genes within a cell comprising: (a) synthesizing one or more siRNA molecules of the invention, which can be chemically-modified, wherein the siRNA strands comprise sequences complementary to RNA of the target genes and wherein the sense strand sequences of the siRNAs comprise sequences identical or
15 substantially similar to the sequences of the target RNAs; and (b) introducing the siRNA molecules into a cell under conditions suitable to modulate the expression of the target genes in the cell.

In another embodiment, the invention features a method for modulating the expression of more than one target gene within a cell comprising: (a) synthesizing a
20 siRNA molecule of the invention, which can be chemically-modified, wherein one of the siRNA strands comprises a sequence complementary to RNA of the target gene and wherein the sense strand sequence of the siRNA comprises a sequence identical or substantially similar to the sequences of the target RNAs; and (b) introducing the siRNA molecule into a cell under conditions suitable to modulate the expression of the target
25 genes in the cell.

In one embodiment, siRNA molecules of the invention are used as reagents in *ex vivo* applications. For example, siRNA reagents are introduced into tissue or cells that are transplanted into a subject for therapeutic effect. The cells and/or tissue can be derived from an organism or subject that later receives the explant, or can be derived from
30 another organism or subject prior to transplantation. The siRNA molecules can be used to

modulate the expression of one or more target genes in the cells or tissue, such that the cells or tissue obtain a desired phenotype or are able to perform a function when transplanted in vivo. In one embodiment, certain target cells from a patient are extracted. These extracted cells are contacted with siNAs targeting a specific nucleotide sequence within the cells under conditions suitable for uptake of the siNAs by these cells (e.g. using delivery reagents such as cationic lipids, liposomes and the like or using techniques such as electroporation to facilitate the delivery of siNAs into cells). The cells are then reintroduced back into the same patient or other patients. In one embodiment, the invention features a method of modulating the expression of a target gene in a tissue explant comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the target gene; and (b) introducing the siNA molecule into a cell of the tissue explant derived from a particular organism under conditions suitable to modulate the expression of the target gene in the tissue explant. In another embodiment, the method further comprises introducing the tissue explant back into the organism the tissue was derived from or into another organism under conditions suitable to modulate the expression of the target gene in that organism.

In one embodiment, the invention features a method of modulating the expression of a target gene in a tissue explant comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the target gene and wherein the sense strand sequence of the siNA comprises a sequence identical or substantially similar to the sequence of the target RNA; and (b) introducing the siNA molecule into a cell of the tissue explant derived from a particular organism under conditions suitable to modulate the expression of the gene in the tissue explant. In another embodiment, the method further comprises introducing the tissue explant back into the organism the tissue was derived from or into another organism under conditions suitable to modulate the expression of the target gene in that organism.

In another embodiment, the invention features a method of modulating the expression of more than one target gene in a tissue explant comprising: (a) synthesizing siNA molecules of the invention, which can be chemically-modified, wherein one of the

siNA strands comprises a sequence complementary to RNA of the target genes; and (b) introducing the siNA molecules into a cell of the tissue explant derived from a particular organism under conditions suitable to modulate the expression of the target genes in the tissue explant. In another embodiment, the method further comprises introducing the
5 tissue explant back into the organism the tissue was derived from or into another organism under conditions suitable to modulate the expression of the target genes in that organism.

In one embodiment, the invention features a method of modulating the expression of a target gene in a subject or organism comprising: (a) synthesizing a siNA molecule
10 of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the target gene; and (b) introducing the siNA molecule into the subject or organism under conditions suitable to modulate the expression of the target gene in the subject or organism. The level of protein or RNA can be determined using various methods well-known in the art.

In another embodiment, the invention features a method of modulating the expression of more than one target gene in a subject or organism comprising: (a) synthesizing siNA molecules of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the target genes; and (b) introducing the siNA molecules into the subject or organism under
20 conditions suitable to modulate the expression of the target genes in the subject or organism. The level of protein or RNA can be determined as is known in the art.

In one embodiment, the invention features a method for modulating the expression of a target gene within a cell comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein the siNA comprises a single
25 stranded sequence having complementarity to RNA of the target gene; and (b) introducing the siNA molecule into a cell under conditions suitable to modulate the expression of the target gene in the cell.

In another embodiment, the invention features a method for modulating the expression of more than one target gene within a cell comprising: (a) synthesizing siNA
30 molecules of the invention, which can be chemically-modified, wherein the siNA

comprises a single stranded sequence having complementarity to RNA of the target gene; and (b) contacting the cell *in vitro* or *in vivo* with the siNA molecule under conditions suitable to modulate the expression of the target genes in the cell.

In one embodiment, the invention features a method of modulating the expression of a target gene in a tissue explant comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein the siNA comprises a single stranded sequence having complementarity to RNA of the gene; and (b) contacting a cell of the tissue explant derived from a particular subject or organism with the siNA molecule under conditions suitable to modulate the expression of the target gene in the tissue explant. In another embodiment, the method further comprises introducing the tissue explant back into the subject or organism the tissue was derived from or into another subject or organism under conditions suitable to modulate the expression of the target gene in that subject or organism.

In another embodiment, the invention features a method of modulating the expression of more than one target gene in a tissue explant comprising: (a) synthesizing siNA molecules of the invention, which can be chemically-modified, wherein the siNA comprises a single stranded sequence having complementarity to RNA of the target gene; and (b) introducing the siNA molecules into a cell of the tissue explant derived from a particular subject or organism under conditions suitable to modulate the expression of the target genes in the tissue explant. In another embodiment, the method further comprises introducing the tissue explant back into the subject or organism the tissue was derived from or into another subject or organism under conditions suitable to modulate the expression of the target genes in that subject or organism.

In one embodiment, the invention features a method of modulating the expression of a target gene in a subject or organism comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein the siNA comprises a single stranded sequence having complementarity to RNA of the target gene; and (b) introducing the siNA molecule into the subject or organism under conditions suitable to modulate the expression of the target gene in the subject or organism.

In another embodiment, the invention features a method of modulating the expression of more than one target gene in a subject or organism comprising: (a) synthesizing siNA molecules of the invention, which can be chemically-modified, wherein the siNA comprises a single stranded sequence having complementarity to RNA of the target gene; and (b) introducing the siNA molecules into the subject or organism under conditions suitable to modulate the expression of the target genes in the subject or organism.

In one embodiment, the invention features a method of modulating the expression of a target gene in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of the target gene in the subject or organism.

In one embodiment, the invention features a method for treating or preventing a disease, disorder, trait or condition related to gene expression in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of the target gene in the subject or organism. The reduction of gene expression and thus reduction in the level of the respective protein/RNA relieves, to some extent, the symptoms of the disease, disorder, trait or condition.

In one embodiment, the invention features a method for treating or preventing cancer in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of a target gene associated with the maintenance or development of cancer in the subject or organism.

In one embodiment, the invention features a method for treating or preventing a proliferative disease, disorder, trait or condition in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of a target gene associated with the maintenance or development of the proliferative disease, disorder, trait or condition in the subject or organism.

In one embodiment, the invention features a method for treating or preventing transplant and/or tissue rejection (allograft rejection) in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of a target gene associated with transplant and/or tissue rejection (allograft rejection) in the subject or organism.

In one embodiment, the invention features a method for treating or preventing an autoimmune disease, disorder, trait or condition in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of a target gene associated with the maintenance or development of the autoimmune disease, disorder, trait or condition in the subject or organism.

In one embodiment, the invention features a method for treating or preventing an infectious disease, disorder, trait or condition in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of a target gene associated with the maintenance or development of the infectious disease, disorder, trait or condition in the subject or organism.

In one embodiment, the invention features a method for treating or preventing an age-related disease, disorder, trait or condition in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of a target gene associated with the maintenance or development of the age-related disease, disorder, trait or condition in the subject or organism.

In one embodiment, the invention features a method for treating or preventing a neurologic or neurodegenerative disease, disorder, trait or condition in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of a target gene associated with the maintenance or development of the neurologic or neurodegenerative disease, disorder, trait or condition in the subject or organism.

In one embodiment, the invention features a method for treating or preventing a metabolic disease, disorder, trait or condition in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of a target gene associated with the maintenance or development of the metabolic disease, disorder, trait or condition in the subject or organism.

In one embodiment, the invention features a method for treating or preventing an cardiovascular disease, disorder, trait or condition in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of a target gene associated with the maintenance or development of the cardiovascular disease, disorder, trait or condition in the subject or organism.

In one embodiment, the invention features a method for treating or preventing a respiratory disease, disorder, trait or condition in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of a target gene associated with the maintenance or development of the respiratory disease, disorder, trait or condition in the subject or organism.

In one embodiment, the invention features a method for treating or preventing an ocular disease, disorder, trait or condition in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of a target gene associated with the maintenance or development of the ocular disease, disorder, trait or condition in the subject or organism.

In one embodiment, the invention features a method for treating or preventing a dermatological disease, disorder, trait or condition in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of a target gene associated with the maintenance or development of the dermatological disease, disorder, trait or condition in the subject or organism.

In one embodiment, the invention features a method for treating or preventing a disease, disorder, trait or condition in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of a target gene associated with the maintenance or development of the disease, disorder, trait or condition in the subject or organism.

In one embodiment, the invention features a method for treating or preventing a liver disease, disorder, trait or condition in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of a target gene associated with the maintenance or development of the liver disease, disorder, trait or condition in the subject or organism.

In one embodiment, the invention features a method for treating or preventing a kidney disease, disorder, trait or condition in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of a target gene associated with the maintenance or development of the kidney disease, disorder, trait or condition in the subject or organism.

In one embodiment, the invention features a method for treating or preventing a bladder disease, disorder, trait or condition in a subject or organism comprising contacting the subject or organism with a siNA molecule of the invention under conditions suitable to modulate the expression of a target gene associated with the maintenance or development of the bladder disease, disorder, trait or condition in the subject or organism.

In another embodiment, the invention features a method of modulating the expression of more than one target gene in a subject or organism comprising contacting the subject or organism with one or more siNA molecules of the invention under conditions suitable to modulate the expression of the genes in the subject or organism.

The siNA molecules of the invention can be designed to down regulate or inhibit target gene expression through RNAi targeting of a variety of RNA molecules. In one embodiment, the siNA molecules of the invention are used to target various RNAs corresponding to a target gene. Non-limiting examples of such RNAs include messenger

RNA (mRNA), alternate RNA splice variants of target gene(s), post-transcriptionally modified RNA of target gene(s), pre-mRNA of target gene(s), and/or RNA templates. If alternate splicing produces a family of transcripts that are distinguished by usage of appropriate exons, the instant invention can be used to inhibit gene expression through the appropriate exons to specifically inhibit or to distinguish among the functions of gene family members. For example, a protein that contains an alternatively spliced transmembrane domain can be expressed in both membrane bound and secreted forms. Use of the invention to target the exon containing the transmembrane domain can be used to determine the functional consequences of pharmaceutical targeting of membrane bound as opposed to the secreted form of the protein. Non-limiting examples of applications of the invention relating to targeting these RNA molecules include therapeutic pharmaceutical applications, pharmaceutical discovery applications, molecular diagnostic and gene function applications, and gene mapping, for example using single nucleotide polymorphism mapping with siRNA molecules of the invention. Such applications can be implemented using known gene sequences or from partial sequences available from an expressed sequence tag (EST).

In another embodiment, the siRNA molecules of the invention are used to target conserved sequences corresponding to a target gene family or target gene families. As such, siRNA molecules targeting multiple gene targets can provide increased therapeutic effect. In addition, siRNA can be used to characterize pathways of gene function in a variety of applications. For example, the present invention can be used to inhibit the activity of target gene(s) in a pathway to determine the function of uncharacterized gene(s) in gene function analysis, mRNA function analysis, or translational analysis. The invention can be used to determine potential target gene pathways involved in various diseases and conditions toward pharmaceutical development. The invention can be used to understand pathways of gene expression involved in diseases, traits, disorders, and/or conditions described herein or otherwise known in the art.

In one embodiment, siRNA molecule(s) and/or methods of the invention are used to down regulate the expression of gene(s) that encode RNA referred to by Genbank Accession, for example, target genes encoding RNA sequence(s) referred to herein by Genbank Accession number, for example, Genbank Accession Nos. shown in Table I.

In one embodiment, the invention features a method comprising: (a) generating a library of siRNA constructs having a predetermined complexity; and (b) assaying the siRNA constructs of (a) above, under conditions suitable to determine RNAi target sites within the target RNA sequence. In one embodiment, the siRNA molecules of (a) have strands of a fixed length, for example, about 23 nucleotides in length. In another embodiment, the siRNA molecules of (a) are of differing length, for example having strands of about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides in length. In one embodiment, the assay can comprise a reconstituted *in vitro* siRNA assay as described herein. In another embodiment, the assay can comprise a cell culture system in which target RNA is expressed. In another embodiment, fragments of target RNA are analyzed for detectable levels of cleavage, for example by gel electrophoresis, northern blot analysis, or RNase protection assays, to determine the most suitable target site(s) within the target RNA sequence. The target RNA sequence can be obtained as is known in the art, for example, by cloning and/or transcription for *in vitro* systems, and by cellular expression in *in vivo* systems.

In one embodiment, the invention features a method comprising: (a) generating a randomized library of siRNA constructs having a predetermined complexity, such as of 4^N , where N represents the number of base paired nucleotides in each of the siRNA construct strands (eg. for a siRNA construct having 21 nucleotide sense and antisense strands with 19 base pairs, the complexity would be 4^{19}); and (b) assaying the siRNA constructs of (a) above, under conditions suitable to determine RNAi target sites within the target RNA sequence. In another embodiment, the siRNA molecules of (a) have strands of a fixed length, for example about 23 nucleotides in length. In yet another embodiment, the siRNA molecules of (a) are of differing length, for example having strands of about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides in length. In one embodiment, the assay can comprise a reconstituted *in vitro* siRNA assay as described in Example 6 herein. In another embodiment, the assay can comprise a cell culture system in which target RNA is expressed. In another embodiment, fragments of target RNA are analyzed for detectable levels of cleavage, for example, by gel electrophoresis, northern blot analysis, or RNase protection assays, to determine the most suitable target site(s) within the target target RNA sequence. The target target

RNA sequence can be obtained as is known in the art, for example, by cloning and/or transcription for *in vitro* systems, and by cellular expression in *in vivo* systems.

In another embodiment, the invention features a method comprising: (a) analyzing the sequence of a RNA target encoded by a target gene; (b) synthesizing one or more sets of siNA molecules having sequence complementary to one or more regions of the RNA of (a); and (c) assaying the siNA molecules of (b) under conditions suitable to determine RNAi targets within the target RNA sequence. In one embodiment, the siNA molecules of (b) have strands of a fixed length, for example about 23 nucleotides in length. In another embodiment, the siNA molecules of (b) are of differing length, for example having strands of about 15 to about 30 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) nucleotides in length. In one embodiment, the assay can comprise a reconstituted *in vitro* siNA assay as described herein. In another embodiment, the assay can comprise a cell culture system in which target RNA is expressed. Fragments of target RNA are analyzed for detectable levels of cleavage, for example by gel electrophoresis, northern blot analysis, or RNase protection assays, to determine the most suitable target site(s) within the target RNA sequence. The target RNA sequence can be obtained as is known in the art, for example, by cloning and/or transcription for *in vitro* systems, and by expression in *in vivo* systems.

By "target site" is meant a sequence within a target RNA that is "targeted" for cleavage mediated by a siNA construct which contains sequences within its antisense region that are complementary to the target sequence.

By "detectable level of cleavage" is meant cleavage of target RNA (and formation of cleaved product RNAs) to an extent sufficient to discern cleavage products above the background of RNAs produced by random degradation of the target RNA. Production of cleavage products from 1-5% of the target RNA is sufficient to detect above the background for most methods of detection.

In one embodiment, the invention features a composition comprising a siNA molecule of the invention, which can be chemically-modified, in a pharmaceutically acceptable carrier or diluent. In another embodiment, the invention features a pharmaceutical composition comprising siNA molecules of the invention, which can be

chemically-modified, targeting one or more genes in a pharmaceutically acceptable carrier or diluent. In another embodiment, the invention features a method for diagnosing a disease or condition in a subject comprising administering to the subject a composition of the invention under conditions suitable for the diagnosis of the disease or condition in the subject. In another embodiment, the invention features a method for treating or preventing a disease or condition in a subject, comprising administering to the subject a composition of the invention under conditions suitable for the treatment or prevention of the disease or condition in the subject, alone or in conjunction with one or more other therapeutic compounds. In yet another embodiment, the invention features a method for treating or preventing diseases, traits, disorders, and/or conditions in a subject or organism comprising administering to the subject a composition of the invention under conditions suitable for the treatment or prevention of the disease, trait, disorder, and/or condition in the subject or organism.

In another embodiment, the invention features a method for validating a gene target, comprising: (a) synthesizing a siRNA molecule of the invention, which can be chemically-modified, wherein one of the siRNA strands includes a sequence complementary to RNA of a target gene; (b) introducing the siRNA molecule into a cell, tissue, subject, or organism under conditions suitable for modulating expression of the target gene in the cell, tissue, subject, or organism; and (c) determining the function of the target gene by assaying for any phenotypic change in the cell, tissue, subject, or organism.

In another embodiment, the invention features a method for validating a gene target comprising: (a) synthesizing a siRNA molecule of the invention, which can be chemically-modified, wherein one of the siRNA strands includes a sequence complementary to RNA of a target gene; (b) introducing the siRNA molecule into a biological system under conditions suitable for modulating expression of the target gene in the biological system; and (c) determining the function of the gene by assaying for any phenotypic change in the biological system.

By "biological system" is meant, material, in a purified or unpurified form, from biological sources, including but not limited to human or animal, wherein the system comprises the components required for RNAi activity. The term "biological system"

includes, for example, a cell, tissue, subject, or organism, or extract thereof. The term biological system also includes reconstituted RNAi systems that can be used in an *in vitro* setting.

By "phenotypic change" is meant any detectable change to a cell that occurs in response to contact or treatment with a nucleic acid molecule of the invention (e.g., siNA). Such detectable changes include, but are not limited to, changes in shape, size, proliferation, motility, protein expression or RNA expression or other physical or chemical changes as can be assayed by methods known in the art. The detectable change can also include expression of reporter genes/molecules such as Green Florescent Protein (GFP) or various tags that are used to identify an expressed protein or any other cellular component that can be assayed.

In one embodiment, the invention features a kit containing a siNA molecule of the invention, which can be chemically-modified, that can be used to modulate the expression of a target gene in a biological system, including, for example, in a cell, tissue, subject, or organism. In another embodiment, the invention features a kit containing more than one siNA molecule of the invention, which can be chemically-modified, that can be used to modulate the expression of more than one target gene in a biological system, including, for example, in a cell, tissue, subject, or organism.

In one embodiment, the invention features a cell containing one or more siNA molecules of the invention, which can be chemically-modified. In another embodiment, the cell containing a siNA molecule of the invention is a mammalian cell. In yet another embodiment, the cell containing a siNA molecule of the invention is a human cell.

In one embodiment, the synthesis of a siNA molecule of the invention, which can be chemically-modified, comprises: (a) synthesis of two complementary strands of the siNA molecule; (b) annealing the two complementary strands together under conditions suitable to obtain a double-stranded siNA molecule. In another embodiment, synthesis of the two complementary strands of the siNA molecule is by solid phase oligonucleotide synthesis. In yet another embodiment, synthesis of the two complementary strands of the siNA molecule is by solid phase tandem oligonucleotide synthesis.

In one embodiment, the invention features a method for synthesizing a siNA duplex molecule comprising: (a) synthesizing a first oligonucleotide sequence strand of the siNA molecule, wherein the first oligonucleotide sequence strand comprises a cleavable linker molecule that can be used as a scaffold for the synthesis of the second oligonucleotide sequence strand of the siNA; (b) synthesizing the second oligonucleotide sequence strand of siNA on the scaffold of the first oligonucleotide sequence strand, wherein the second oligonucleotide sequence strand further comprises a chemical moiety than can be used to purify the siNA duplex; (c) cleaving the linker molecule of (a) under conditions suitable for the two siNA oligonucleotide strands to hybridize and form a stable duplex; and (d) purifying the siNA duplex utilizing the chemical moiety of the second oligonucleotide sequence strand. In one embodiment, cleavage of the linker molecule in (c) above takes place during deprotection of the oligonucleotide, for example, under hydrolysis conditions using an alkylamine base such as methylamine. In one embodiment, the method of synthesis comprises solid phase synthesis on a solid support such as controlled pore glass (CPG) or polystyrene, wherein the first sequence of (a) is synthesized on a cleavable linker, such as a succinyl linker, using the solid support as a scaffold. The cleavable linker in (a) used as a scaffold for synthesizing the second strand can comprise similar reactivity as the solid support derivatized linker, such that cleavage of the solid support derivatized linker and the cleavable linker of (a) takes place concomitantly. In another embodiment, the chemical moiety of (b) that can be used to isolate the attached oligonucleotide sequence comprises a trityl group, for example a dimethoxytrityl group, which can be employed in a trityl-on synthesis strategy as described herein. In yet another embodiment, the chemical moiety, such as a dimethoxytrityl group, is removed during purification, for example, using acidic conditions.

In a further embodiment, the method for siNA synthesis is a solution phase synthesis or hybrid phase synthesis wherein both strands of the siNA duplex are synthesized in tandem using a cleavable linker attached to the first sequence which acts a scaffold for synthesis of the second sequence. Cleavage of the linker under conditions suitable for hybridization of the separate siNA sequence strands results in formation of the double-stranded siNA molecule.

In another embodiment, the invention features a method for synthesizing a siNA duplex molecule comprising: (a) synthesizing one oligonucleotide sequence strand of the siNA molecule, wherein the sequence comprises a cleavable linker molecule that can be used as a scaffold for the synthesis of another oligonucleotide sequence; (b) synthesizing a second oligonucleotide sequence having complementarity to the first sequence strand on the scaffold of (a), wherein the second sequence comprises the other strand of the double-stranded siNA molecule and wherein the second sequence further comprises a chemical moiety that can be used to isolate the attached oligonucleotide sequence; (c) purifying the product of (b) utilizing the chemical moiety of the second oligonucleotide sequence strand under conditions suitable for isolating the full-length sequence comprising both siNA oligonucleotide strands connected by the cleavable linker and under conditions suitable for the two siNA oligonucleotide strands to hybridize and form a stable duplex. In one embodiment, cleavage of the linker molecule in (c) above takes place during deprotection of the oligonucleotide, for example, under hydrolysis conditions. In another embodiment, cleavage of the linker molecule in (c) above takes place after deprotection of the oligonucleotide. In another embodiment, the method of synthesis comprises solid phase synthesis on a solid support such as controlled pore glass (CPG) or polystyrene, wherein the first sequence of (a) is synthesized on a cleavable linker, such as a succinyl linker, using the solid support as a scaffold. The cleavable linker in (a) used as a scaffold for synthesizing the second strand can comprise similar reactivity or differing reactivity as the solid support derivatized linker, such that cleavage of the solid support derivatized linker and the cleavable linker of (a) takes place either concomitantly or sequentially. In one embodiment, the chemical moiety of (b) that can be used to isolate the attached oligonucleotide sequence comprises a trityl group, for example a dimethoxytrityl group.

In another embodiment, the invention features a method for making a double-stranded siNA molecule in a single synthetic process comprising: (a) synthesizing an oligonucleotide having a first and a second sequence, wherein the first sequence is complementary to the second sequence, and the first oligonucleotide sequence is linked to the second sequence via a cleavable linker, and wherein a terminal 5'-protecting group, for example, a 5'-O-dimethoxytrityl group (5'-O-DMT) remains on the oligonucleotide having the second sequence; (b) deprotecting the oligonucleotide whereby the

deprotection results in the cleavage of the linker joining the two oligonucleotide sequences; and (c) purifying the product of (b) under conditions suitable for isolating the double-stranded siNA molecule, for example using a trityl-on synthesis strategy as described herein.

- 5 In another embodiment, the method of synthesis of siNA molecules of the invention comprises the teachings of Scaringe *et al.*, US Patent Nos. 5,889,136; 6,008,400; and 6,111,086, incorporated by reference herein in their entirety.

- 10 In one embodiment, the invention features siNA constructs that mediate RNAi against a target polynucleotide (e.g., DNA or RNA), wherein the siNA construct comprises one or more chemical modifications, for example, one or more chemical modifications having any of Formulae I-VII or any combination thereof that increases the nuclease resistance of the siNA construct.

- 15 In another embodiment, the invention features a method for generating siNA molecules with increased nuclease resistance comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having increased nuclease resistance.

- 20 In another embodiment, the invention features a method for generating siNA molecules with improved toxicologic profiles (e.g., have attenuated or no immunostimulatory properties) comprising (a) introducing nucleotides having any of Formula I-VII (e.g., siNA motifs referred to in Table II) or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved toxicologic profiles.

- 25 In another embodiment, the invention features a method for generating siNA molecules that do not stimulate an interferon response (e.g., no interferon response or attenuated interferon response) in a cell, subject, or organism, comprising (a) introducing nucleotides having any of Formula I-VII (e.g., siNA motifs referred to in Table II) or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules that do not stimulate an interferon response.
- 30

By "improved toxicologic profile", is meant that the chemically modified siNA construct exhibits decreased toxicity in a cell, subject, or organism compared to an unmodified siNA or siNA molecule having fewer modifications or modifications that are less effective in imparting improved toxicology. In a non-limiting example, siNA molecules with improved toxicologic profiles are associated with a decreased or attenuated immunostimulatory response in a cell, subject, or organism compared to an unmodified siNA or siNA molecule having fewer modifications or modifications that are less effective in imparting improved toxicology. In one embodiment, a siNA molecule with an improved toxicological profile comprises no ribonucleotides. In one embodiment, a siNA molecule with an improved toxicological profile comprises less than 5 ribonucleotides (e.g., 1, 2, 3, or 4 ribonucleotides). In one embodiment, a siNA molecule with an improved toxicological profile comprises Stab 7, Stab 8, Stab 11, Stab 12, Stab 13, Stab 16, Stab 17, Stab 18, Stab 19, Stab 20, Stab 23, Stab 24, Stab 25, Stab 26, Stab 27, Stab 28, Stab 29, Stab 30, Stab 31, Stab 32 or any combination thereof (see Table II). In one embodiment, the level of immunostimulatory response associated with a given siNA molecule can be measured as is known in the art, for example by determining the level of PKR/interferon response, proliferation, B-cell activation, and/or cytokine production in assays to quantitate the immunostimulatory response of particular siNA molecules (see, for example, Leifer *et al.*, 2003, *J Immunother.* 26, 313-9; and U.S. Patent No. 5968909, incorporated in its entirety by reference).

In one embodiment, the invention features siNA constructs that mediate RNAi against a target polynucleotide (e.g., DNA or RNA), wherein the siNA construct comprises one or more chemical modifications described herein that modulates the binding affinity between the sense and antisense strands of the siNA construct.

In another embodiment, the invention features a method for generating siNA molecules with increased binding affinity between the sense and antisense strands of the siNA molecule comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having increased binding affinity between the sense and antisense strands of the siNA molecule.

In one embodiment, the invention features siNA constructs that mediate RNAi against a target polynucleotide (e.g., DNA or RNA), wherein the siNA construct comprises one or more chemical modifications described herein that modulates the binding affinity between the antisense strand of the siNA construct and a complementary target RNA sequence within a cell.

In one embodiment, the invention features siNA constructs that mediate RNAi against a target polynucleotide (e.g., DNA or RNA), wherein the siNA construct comprises one or more chemical modifications described herein that modulates the binding affinity between the antisense strand of the siNA construct and a complementary target DNA sequence within a cell.

In another embodiment, the invention features a method for generating siNA molecules with increased binding affinity between the antisense strand of the siNA molecule and a complementary target RNA sequence comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having increased binding affinity between the antisense strand of the siNA molecule and a complementary target RNA sequence.

In another embodiment, the invention features a method for generating siNA molecules with increased binding affinity between the antisense strand of the siNA molecule and a complementary target DNA sequence comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having increased binding affinity between the antisense strand of the siNA molecule and a complementary target DNA sequence.

In one embodiment, the invention features siNA constructs that mediate RNAi against a target polynucleotide (e.g., DNA or RNA), wherein the siNA construct comprises one or more chemical modifications described herein that modulate the polymerase activity of a cellular polymerase capable of generating additional endogenous siNA molecules having sequence homology to the chemically-modified siNA construct.

In another embodiment, the invention features a method for generating siNA molecules capable of mediating increased polymerase activity of a cellular polymerase capable of generating additional endogenous siNA molecules having sequence homology to a chemically-modified siNA molecule comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules capable of mediating increased polymerase activity of a cellular polymerase capable of generating additional endogenous siNA molecules having sequence homology to the chemically-modified siNA molecule.

10 In one embodiment, the invention features chemically-modified siNA constructs that mediate RNAi against a target polynucleotide (e.g., DNA or RNA) in a cell, wherein the chemical modifications do not significantly effect the interaction of siNA with a target RNA molecule, DNA molecule and/or proteins or other factors that are essential for RNAi in a manner that would decrease the efficacy of RNAi mediated by such siNA constructs.

15 In another embodiment, the invention features a method for generating siNA molecules with improved RNAi activity against a target polynucleotide (e.g., DNA or RNA) comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved RNAi activity.

20 In yet another embodiment, the invention features a method for generating siNA molecules with improved RNAi activity against a target RNA comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved RNAi activity against the target RNA.

In yet another embodiment, the invention features a method for generating siNA molecules with improved RNAi activity against a target DNA comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA

molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved RNAi activity against the target DNA.

In one embodiment, the invention features siNA constructs that mediate RNAi against a target polynucleotide (e.g., DNA or RNA), wherein the siNA construct
5 comprises one or more chemical modifications described herein that modulates the cellular uptake of the siNA construct.

In another embodiment, the invention features a method for generating siNA molecules against a target polynucleotide (e.g., DNA or RNA) with improved cellular uptake comprising (a) introducing nucleotides having any of Formula I-VII or any
10 combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved cellular uptake.

In one embodiment, the invention features siNA constructs that mediate RNAi against a target polynucleotide (e.g., DNA or RNA), wherein the siNA construct
15 comprises one or more chemical modifications described herein that increases the bioavailability of the siNA construct, for example, by attaching polymeric conjugates such as polyethyleneglycol or equivalent conjugates that improve the pharmacokinetics of the siNA construct, or by attaching conjugates that target specific tissue types or cell types *in vivo*. Non-limiting examples of such conjugates are described in Vargeese *et al.*,
20 U.S. Serial No. 10/201,394 incorporated by reference herein.

In one embodiment, the invention features a method for generating siNA molecules of the invention with improved bioavailability comprising (a) introducing a conjugate into the structure of a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved
25 bioavailability. Such conjugates can include ligands for cellular receptors, such as peptides derived from naturally occurring protein ligands; protein localization sequences, including cellular ZIP code sequences; antibodies; nucleic acid aptamers; vitamins and other co-factors, such as folate and N-acetylgalactosamine; polymers, such as polyethyleneglycol (PEG); phospholipids; cholesterol; polyamines, such as spermine or
30 spermidine; and others.

In one embodiment, the invention features a double stranded short interfering nucleic acid (siNA) molecule that comprises a first nucleotide sequence complementary to a target RNA sequence or a portion thereof, and a second sequence having complementarity to said first sequence, wherein said second sequence is chemically
5 modified in a manner that it can no longer act as a guide sequence for efficiently mediating RNA interference and/or be recognized by cellular proteins that facilitate RNAi.

In one embodiment, the invention features a double stranded short interfering nucleic acid (siNA) molecule that comprises a first nucleotide sequence complementary
10 to a target RNA sequence or a portion thereof, and a second sequence having complementarity to said first sequence, wherein the second sequence is designed or modified in a manner that prevents its entry into the RNAi pathway as a guide sequence or as a sequence that is complementary to a target nucleic acid (e.g., RNA) sequence. Such design or modifications are expected to enhance the activity of siNA and/or
15 improve the specificity of siNA molecules of the invention. These modifications are also expected to minimize any off-target effects and/or associated toxicity.

In one embodiment, the invention features a double stranded short interfering nucleic acid (siNA) molecule that comprises a first nucleotide sequence complementary to a target RNA sequence or a portion thereof, and a second sequence having
20 complementarity to said first sequence, wherein said second sequence is incapable of acting as a guide sequence for mediating RNA interference.

In one embodiment, the invention features a double stranded short interfering nucleic acid (siNA) molecule that comprises a first nucleotide sequence complementary to a target RNA sequence or a portion thereof, and a second sequence having
25 complementarity to said first sequence, wherein said second sequence does not have a terminal 5'-hydroxyl (5'-OH) or 5'-phosphate group.

In one embodiment, the invention features a double stranded short interfering nucleic acid (siNA) molecule that comprises a first nucleotide sequence complementary to a target RNA sequence or a portion thereof, and a second sequence having
30 complementarity to said first sequence, wherein said second sequence comprises a

terminal cap moiety at the 5'-end of said second sequence. In one embodiment, the terminal cap moiety comprises an inverted abasic, inverted deoxy abasic, inverted nucleotide moiety, a group shown in **Figure 10**, an alkyl or cycloalkyl group, a heterocycle, or any other group that prevents RNAi activity in which the second sequence serves as a guide sequence or template for RNAi.

In one embodiment, the invention features a double stranded short interfering nucleic acid (siNA) molecule that comprises a first nucleotide sequence complementary to a target RNA sequence or a portion thereof, and a second sequence having complementarity to said first sequence, wherein said second sequence comprises a terminal cap moiety at the 5'-end and 3'-end of said second sequence. In one embodiment, each terminal cap moiety individually comprises an inverted abasic, inverted deoxy abasic, inverted nucleotide moiety, a group shown in **Figure 10**, an alkyl or cycloalkyl group, a heterocycle, or any other group that prevents RNAi activity in which the second sequence serves as a guide sequence or template for RNAi.

In one embodiment, the invention features a method for generating siNA molecules of the invention with improved specificity for down regulating or inhibiting the expression of a target nucleic acid (e.g., a DNA or RNA such as a gene or its corresponding RNA), comprising (a) introducing one or more chemical modifications into the structure of a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved specificity. In another embodiment, the chemical modification used to improve specificity comprises terminal cap modifications at the 5'-end, 3'-end, or both 5' and 3'-ends of the siNA molecule. The terminal cap modifications can comprise, for example, structures shown in **Figure 10** (e.g. inverted deoxyabasic moieties) or any other chemical modification that renders a portion of the siNA molecule (e.g. the sense strand) incapable of mediating RNA interference against an off target nucleic acid sequence. In a non-limiting example, a siNA molecule is designed such that only the antisense sequence of the siNA molecule can serve as a guide sequence for RISC mediated degradation of a corresponding target RNA sequence. This can be accomplished by rendering the sense sequence of the siNA inactive by introducing chemical modifications to the sense strand that preclude recognition of the sense strand as a guide sequence by RNAi machinery. In one

embodiment, such chemical modifications comprise any chemical group at the 5'-end or the sense strand of the siNA, or any other group that serves to render the sense strand inactive as a guide sequence for mediating RNA interference. These modifications, for example, can result in a molecule where the 5'-end of the sense strand no longer has a free 5'-hydroxyl (5'-OH) or a free 5'-phosphate group (e.g., phosphate, diphosphate, triphosphate, cyclic phosphate etc.). Non-limiting examples of such siNA constructs are described herein, such as "Stab 9/10", "Stab 7/8", "Stab 7/19", "Stab 17/22", "Stab 23/24", "Stab 24/25", and "Stab 24/26" (e.g., any siNA having Stab 7, 9, 17, 23, or 24 sense strands) chemistries and variants thereof (see Table II) wherein the 5'-end and 3'-end of the sense strand of the siNA do not comprise a hydroxyl group or phosphate group.

In one embodiment, the invention features a method for generating siNA molecules of the invention with improved specificity for down regulating or inhibiting the expression of a target nucleic acid (e.g., a DNA or RNA such as a gene or its corresponding RNA), comprising introducing one or more chemical modifications into the structure of a siNA molecule that prevent a strand or portion of the siNA molecule from acting as a template or guide sequence for RNAi activity. In one embodiment, the inactive strand or sense region of the siNA molecule is the sense strand or sense region of the siNA molecule, i.e. the strand or region of the siNA that does not have complementarity to the target nucleic acid sequence. In one embodiment, such chemical modifications comprise any chemical group at the 5'-end of the sense strand or region of the siNA that does not comprise a 5'-hydroxyl (5'-OH) or 5'-phosphate group, or any other group that serves to render the sense strand or sense region inactive as a guide sequence for mediating RNA interference. Non-limiting examples of such siNA constructs are described herein, such as "Stab 9/10", "Stab 7/8", "Stab 7/19", "Stab 17/22", "Stab 23/24", "Stab 24/25", and "Stab 24/26" (e.g., any siNA having Stab 7, 9, 17, 23, or 24 sense strands) chemistries and variants thereof (see Table II) wherein the 5'-end and 3'-end of the sense strand of the siNA do not comprise a hydroxyl group or phosphate group.

In one embodiment, the invention features a method for screening siNA molecules that are active in mediating RNA interference against a target nucleic acid sequence

comprising (a) generating a plurality of unmodified siNA molecules, (b) screening the siNA molecules of step (a) under conditions suitable for isolating siNA molecules that are active in mediating RNA interference against the target nucleic acid sequence, and (c) introducing chemical modifications (e.g. chemical modifications as described herein or as otherwise known in the art) into the active siNA molecules of (b). In one embodiment, the method further comprises re-screening the chemically modified siNA molecules of step (c) under conditions suitable for isolating chemically modified siNA molecules that are active in mediating RNA interference against the target nucleic acid sequence.

10 In one embodiment, the invention features a method for screening chemically modified siNA molecules that are active in mediating RNA interference against a target nucleic acid sequence comprising (a) generating a plurality of chemically modified siNA molecules (e.g. siNA molecules as described herein or as otherwise known in the art), and (b) screening the siNA molecules of step (a) under conditions suitable for isolating
15 chemically modified siNA molecules that are active in mediating RNA interference against the target nucleic acid sequence.

The term "ligand" refers to any compound or molecule, such as a drug, peptide, hormone, or neurotransmitter, that is capable of interacting with another compound, such as a receptor, either directly or indirectly. The receptor that interacts with a ligand can be
20 present on the surface of a cell or can alternately be an intercellular receptor. Interaction of the ligand with the receptor can result in a biochemical reaction, or can simply be a physical interaction or association.

In another embodiment, the invention features a method for generating siNA molecules of the invention with improved bioavailability comprising (a) introducing an
25 excipient formulation to a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved bioavailability. Such excipients include polymers such as cyclodextrins, lipids, cationic lipids, polyamines, phospholipids, nanoparticles, receptors, ligands, and others.

In another embodiment, the invention features a method for generating siNA
30 molecules of the invention with improved bioavailability comprising (a) introducing

nucleotides having any of Formulae I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved bioavailability.

In another embodiment, polyethylene glycol (PEG) can be covalently attached to siNA compounds of the present invention. The attached PEG can be any molecular weight, preferably from about 2,000 to about 50,000 daltons (Da).

The present invention can be used alone or as a component of a kit having at least one of the reagents necessary to carry out the *in vitro* or *in vivo* introduction of RNA to test samples and/or subjects. For example, preferred components of the kit include a siNA molecule of the invention and a vehicle that promotes introduction of the siNA into cells of interest as described herein (e.g., using lipids and other methods of transfection known in the art, see for example Beigelman *et al.*, US 6,395,713). The kit can be used for target validation, such as in determining gene function and/or activity, or in drug optimization, and in drug discovery (see for example Usman *et al.*, USSN 60/402,996).

Such a kit can also include instructions to allow a user of the kit to practice the invention.

The term "short interfering nucleic acid", "siNA", "short interfering RNA", "siRNA", "short interfering nucleic acid molecule", "short interfering oligonucleotide molecule", or "chemically-modified short interfering nucleic acid molecule" as used herein refers to any nucleic acid molecule capable of inhibiting or down regulating gene expression or viral replication, for example by mediating RNA interference "RNAi" or gene silencing in a sequence-specific manner; see for example Zamore *et al.*, 2000, *Cell*, 101, 25-33; Bass, 2001, *Nature*, 411, 428-429; Elbashir *et al.*, 2001, *Nature*, 411, 494-498; and Kreutzer *et al.*, International PCT Publication No. WO 00/44895; Zernicka-Goetz *et al.*, International PCT Publication No. WO 01/36646; Fire, International PCT Publication No. WO 99/32619; Plactinck *et al.*, International PCT Publication No. WO 00/01846; Mello and Fire, International PCT Publication No. WO 01/29058; Deschamps-Depaillette, International PCT Publication No. WO 99/07409; and Li *et al.*, International PCT Publication No. WO 00/44914; Allshire, 2002, *Science*, 297, 1818-1819; Volpe *et al.*, 2002, *Science*, 297, 1833-1837; Jenuwein, 2002, *Science*, 297, 2215-2218; and Hall *et al.*, 2002, *Science*, 297, 2232-2237; Hutvagner and Zamore, 2002, *Science*, 297, 2056-60; McManus *et al.*, 2002, *RNA*, 8, 842-850; Reinhart *et al.*, 2002,

Gene & Dev., 16, 1616-1626; and Reinhart & Bartel, 2002, *Science*, 297, 1831). Non limiting examples of siNA molecules of the invention are shown in Figures 4-6 herein. For example the siNA can be a double-stranded polynucleotide molecule comprising self-complementary sense and antisense regions, wherein the antisense region comprises

5 nucleotide sequence that is complementary to nucleotide sequence in a target nucleic acid molecule or a portion thereof and the sense region having nucleotide sequence corresponding to the target nucleic acid sequence or a portion thereof. The siNA can be assembled from two separate oligonucleotides, where one strand is the sense strand and the other is the antisense strand, wherein the antisense and sense strands are self-

10 complementary (i.e. each strand comprises nucleotide sequence that is complementary to nucleotide sequence in the other strand; such as where the antisense strand and sense strand form a duplex or double stranded structure, for example wherein the double stranded region is about 15 to about 30, e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29 or 30 base pairs; the antisense strand comprises nucleotide sequence

15 that is complementary to nucleotide sequence in a target nucleic acid molecule or a portion thereof and the sense strand comprises nucleotide sequence corresponding to the target nucleic acid sequence or a portion thereof (e.g., about 15 to about 25 or more nucleotides of the siNA molecule are complementary to the target nucleic acid or a portion thereof). Alternatively, the siNA is assembled from a single oligonucleotide,

20 where the self-complementary sense and antisense regions of the siNA are linked by means of a nucleic acid based or non-nucleic acid-based linker(s). The siNA can be a polynucleotide with a duplex, asymmetric duplex, hairpin or asymmetric hairpin secondary structure, having self-complementary sense and antisense regions, wherein the antisense region comprises nucleotide sequence that is complementary to nucleotide

25 sequence in a separate target nucleic acid molecule or a portion thereof and the sense region having nucleotide sequence corresponding to the target nucleic acid sequence or a portion thereof. The siNA can be a circular single-stranded polynucleotide having two or more loop structures and a stem comprising self-complementary sense and antisense regions, wherein the antisense region comprises nucleotide sequence that is

30 complementary to nucleotide sequence in a target nucleic acid molecule or a portion thereof and the sense region having nucleotide sequence corresponding to the target nucleic acid sequence or a portion thereof, and wherein the circular polynucleotide can

be processed either *in vivo* or *in vitro* to generate an active siNA molecule capable of mediating RNAi. The siNA can also comprise a single stranded polynucleotide having nucleotide sequence complementary to nucleotide sequence in a target nucleic acid molecule or a portion thereof (for example, where such siNA molecule does not require the presence within the siNA molecule of nucleotide sequence corresponding to the target nucleic acid sequence or a portion thereof), wherein the single stranded polynucleotide can further comprise a terminal phosphate group, such as a 5'-phosphate (see for example Martinez *et al.*, 2002, *Cell*, 110, 563-574 and Schwarz *et al.*, 2002, *Molecular Cell*, 10, 537-568), or 5',3'-diphosphate. In certain embodiments, the siNA molecule of the invention comprises separate sense and antisense sequences or regions, wherein the sense and antisense regions are covalently linked by nucleotide or non-nucleotide linker molecules as is known in the art, or are alternately non-covalently linked by ionic interactions, hydrogen bonding, van der Waals interactions, hydrophobic interactions, and/or stacking interactions. In certain embodiments, the siNA molecules of the invention comprise nucleotide sequence that is complementary to nucleotide sequence of a target gene. In another embodiment, the siNA molecule of the invention interacts with nucleotide sequence of a target gene in a manner that causes inhibition of expression of the target gene. As used herein, siNA molecules need not be limited to those molecules containing only RNA, but further encompasses chemically-modified nucleotides and non-nucleotides. In certain embodiments, the short interfering nucleic acid molecules of the invention lack 2'-hydroxy (2'-OH) containing nucleotides. Applicant describes in certain embodiments short interfering nucleic acids that do not require the presence of nucleotides having a 2'-hydroxy group for mediating RNAi and as such, short interfering nucleic acid molecules of the invention optionally do not include any ribonucleotides (e.g., nucleotides having a 2'-OH group). Such siNA molecules that do not require the presence of ribonucleotides within the siNA molecule to support RNAi can however have an attached linker or linkers or other attached or associated groups, moieties, or chains containing one or more nucleotides with 2'-OH groups. Optionally, siNA molecules can comprise ribonucleotides at about 5, 10, 20, 30, 40, or 50% of the nucleotide positions. The modified short interfering nucleic acid molecules of the invention can also be referred to as short interfering modified oligonucleotides "siMON." As used herein, the term siNA is meant to be equivalent to

other terms used to describe nucleic acid molecules that are capable of mediating sequence specific RNAi, for example short interfering RNA (siRNA), double-stranded RNA (dsRNA), micro-RNA (miRNA), short hairpin RNA (shRNA), short interfering oligonucleotide, short interfering nucleic acid, short interfering modified oligonucleotide, 5 chemically-modified siRNA, post-transcriptional gene silencing RNA (ptgsRNA), and others. In addition, as used herein, the term RNAi is meant to be equivalent to other terms used to describe sequence specific RNA interference, such as post transcriptional gene silencing, translational inhibition, or epigenetics. For example, siNA molecules of the invention can be used to epigenetically silence genes at both the post-transcriptional 10 level or the pre-transcriptional level. In a non-limiting example, epigenetic regulation of gene expression by siNA molecules of the invention can result from siNA mediated modification of chromatin structure or methylation pattern to alter gene expression (see, for example, Verdel *et al.*, 2004, *Science*, 303, 672-676; Pal-Bhadra *et al.*, 2004, *Science*, 303, 669-672; Allshire, 2002, *Science*, 297, 1818-1819; Volpe *et al.*, 2002, *Science*, 297, 1833-1837; Jenuwein, 2002, *Science*, 297, 2215-2218; and Hall *et al.*, 2002, *Science*, 297, 2232-2237).

In one embodiment, a siNA molecule of the invention is a duplex forming oligonucleotide "DFO", (see for example **Figures 14-15** and Vaish *et al.*, USSN 10/727,780 filed December 3, 2003 and International PCT Application No. US04/16390, 20 filed May 24, 2004).

In one embodiment, a siNA molecule of the invention is a multifunctional siNA, (see for example **Figures 16-21** and Jadhav *et al.*, USSN 60/543,480 filed February 10, 2004 and International PCT Application No. US04/16390, filed May 24, 2004). The multifunctional siNA of the invention can comprise sequence targeting, for example, two 25 regions of target RNA.

By "asymmetric hairpin" as used herein is meant a linear siNA molecule comprising an antisense region, a loop portion that can comprise nucleotides or non-nucleotides, and a sense region that comprises fewer nucleotides than the antisense region to the extent that the sense region has enough complementary nucleotides to base pair with the antisense region and form a duplex with loop. For example, an asymmetric hairpin siNA molecule of the invention can comprise an antisense region having length 30

sufficient to mediate RNAi in a cell or in vitro system (e.g. about 15 to about 30, or about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30 nucleotides) and a loop region comprising about 4 to about 12 (e.g., about 4, 5, 6, 7, 8, 9, 10, 11, or 12) nucleotides, and a sense region having about 3 to about 25 (e.g., about 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, or 25) nucleotides that are complementary to the antisense region. The asymmetric hairpin siNA molecule can also comprise a 5'-terminal phosphate group that can be chemically modified. The loop portion of the asymmetric hairpin siNA molecule can comprise nucleotides, non-nucleotides, linker molecules, or conjugate molecules as described herein.

By "asymmetric duplex" as used herein is meant a siNA molecule having two separate strands comprising a sense region and an antisense region, wherein the sense region comprises fewer nucleotides than the antisense region to the extent that the sense region has enough complementary nucleotides to base pair with the antisense region and form a duplex. For example, an asymmetric duplex siNA molecule of the invention can comprise an antisense region having length sufficient to mediate RNAi in a cell or in vitro system (e.g. about 15 to about 30, or about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30 nucleotides) and a sense region having about 3 to about 25 (e.g., about 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, or 25) nucleotides that are complementary to the antisense region.

By "modulate" is meant that the expression of the gene, or level of RNA molecule or equivalent RNA molecules encoding one or more proteins or protein subunits, or activity of one or more proteins or protein subunits is up regulated or down regulated, such that expression, level, or activity is greater than or less than that observed in the absence of the modulator. For example, the term "modulate" can mean "inhibit," but the use of the word "modulate" is not limited to this definition.

By "inhibit", "down-regulate", or "reduce", it is meant that the expression of the gene, or level of RNA molecules or equivalent RNA molecules encoding one or more proteins or protein subunits, or activity of one or more proteins or protein subunits, is reduced below that observed in the absence of the nucleic acid molecules (e.g., siNA) of the invention. In one embodiment, inhibition, down-regulation or reduction with an siNA molecule is below that level observed in the presence of an inactive or attenuated

molecule. In another embodiment, inhibition, down-regulation, or reduction with siRNA molecules is below that level observed in the presence of, for example, an siRNA molecule with scrambled sequence or with mismatches. In another embodiment, inhibition, down-regulation, or reduction of gene expression with a nucleic acid molecule of the instant invention is greater in the presence of the nucleic acid molecule than in its absence. In one embodiment, inhibition, down regulation, or reduction of gene expression is associated with post transcriptional silencing, such as RNAi mediated cleavage of a target nucleic acid molecule (e.g. RNA) or inhibition of translation. In one embodiment, inhibition, down regulation, or reduction of gene expression is associated with pretranscriptional silencing.

By "gene", or "target gene", is meant a nucleic acid that encodes an RNA, for example, nucleic acid sequences including, but not limited to, structural genes encoding a polypeptide. A gene or target gene can also encode a functional RNA (fRNA) or non-coding RNA (ncRNA), such as small temporal RNA (stRNA), micro RNA (miRNA), small nuclear RNA (snRNA), short interfering RNA (siRNA), small nucleolar RNA (snoRNA), ribosomal RNA (rRNA), transfer RNA (tRNA) and precursor RNAs thereof. Such non-coding RNAs can serve as target nucleic acid molecules for siRNA mediated RNA interference in modulating the activity of fRNA or ncRNA involved in functional or regulatory cellular processes. Abberant fRNA or ncRNA activity leading to disease can therefore be modulated by siRNA molecules of the invention. siRNA molecules targeting fRNA and ncRNA can also be used to manipulate or alter the genotype or phenotype of a subject, organism or cell, by intervening in cellular processes such as genetic imprinting, transcription, translation, or nucleic acid processing (e.g., transamination, methylation etc.). The target gene can be a gene derived from a cell, an endogenous gene, a transgene, or exogenous genes such as genes of a pathogen, for example a virus, which is present in the cell after infection thereof. The cell containing the target gene can be derived from or contained in any organism, for example a plant, animal, protozoan, virus, bacterium, or fungus. Non-limiting examples of plants include monocots, dicots, or gymnosperms. Non-limiting examples of animals include vertebrates or invertebrates. Non-limiting examples of fungi include molds or yeasts. For a review, see for example Snyder and Gerstein, 2003, *Science*, 300, 258-260.

By "non-canonical base pair" is meant any non-Watson Crick base pair, such as mismatches and/or wobble base pairs, including flipped mismatches, single hydrogen bond mismatches, trans-type mismatches, triple base interactions, and quadruple base interactions. Non-limiting examples of such non-canonical base pairs include, but are not limited to, AC reverse Hoogsteen, AC wobble, AU reverse Hoogsteen, GU wobble, AA N7 amino, CC 2-carbonyl-amino(H1)-N3-amino(H2), GA sheared, UC 4-carbonyl-amino, UU imino-carbonyl, AC reverse wobble, AU Hoogsteen, AU reverse Watson Crick, CG reverse Watson Crick, GC N3-amino-amino N3, AA N1-amino symmetric, AA N7-amino symmetric, GA N7-N1 amino-carbonyl, GA+ carbonyl-amino N7-N1, GG N1-carbonyl symmetric, GG N3-amino symmetric, CC carbonyl-amino symmetric, CC N3-amino symmetric, UU 2-carbonyl-imino symmetric, UU 4-carbonyl-imino symmetric, AA amino-N3, AA N1-amino, AC amino 2-carbonyl, AC N3-amino, AC N7-amino, AU amino-4-carbonyl, AU N1-imino, AU N3-imino, AU N7-imino, CC carbonyl-amino, GA amino-N1, GA amino-N7, GA carbonyl-amino, GA N3-amino, GC amino-N3, GC carbonyl-amino, GC N3-amino, GC N7-amino, GG amino-N7, GG carbonyl-imino, GG N7-amino, GU amino-2-carbonyl, GU carbonyl-imino, GU imino-2-carbonyl, GU N7-imino, psiU imino-2-carbonyl, UC 4-carbonyl-amino, UC imino-carbonyl, UU imino-4-carbonyl, AC C2-H-N3, GA carbonyl-C2-H, UU imino-4-carbonyl 2 carbonyl-C5-H, AC amino(A) N3(C)-carbonyl, GC imino amino-carbonyl, Gpsi imino-2-carbonyl amino-2- carbonyl, and GU imino amino-2-carbonyl base pairs.

By "target" as used herein is meant, any target protein, peptide, or polypeptide, such as encoded by Genbank Accession Nos. shown in Table I. The term "target" also refers to nucleic acid sequences encoding any protein, peptide, or polypeptide (e.g., DNA and RNA). The term "target" is also meant to include other target encoding sequences, such as other isoforms, mutations, splice variants, and polymorphisms associated with a given target.

By "homologous sequence" is meant, a nucleotide sequence that is shared by one or more polynucleotide sequences, such as genes, gene transcripts and/or non-coding polynucleotides. For example, a homologous sequence can be a nucleotide sequence that is shared by two or more genes encoding related but different proteins, such as different members of a gene family, different protein epitopes, different protein isoforms or

completely divergent genes, such as a cytokine and its corresponding receptors. A homologous sequence can be a nucleotide sequence that is shared by two or more non-coding polynucleotides, such as noncoding DNA or RNA, regulatory sequences, introns, and sites of transcriptional control or regulation. Homologous sequences can also include conserved sequence regions shared by more than one polynucleotide sequence. Homology does not need to be perfect homology (e.g., 100%), as partially homologous sequences are also contemplated by the instant invention (e.g., 99%, 98%, 97%, 96%, 95%, 94%, 93%, 92%, 91%, 90%, 89%, 88%, 87%, 86%, 85%, 84%, 83%, 82%, 81%, 80% etc.).

By "conserved sequence region" is meant, a nucleotide sequence of one or more regions in a polynucleotide does not vary significantly between generations or from one biological system, subject, or organism to another biological system, subject, or organism. The polynucleotide can include both coding and non-coding DNA and RNA.

By "sense region" is meant a nucleotide sequence of a siNA molecule having complementarity to an antisense region of the siNA molecule. In addition, the sense region of a siNA molecule can comprise a nucleic acid sequence having homology with a target nucleic acid sequence.

By "antisense region" is meant a nucleotide sequence of a siNA molecule having complementarity to a target nucleic acid sequence. In addition, the antisense region of a siNA molecule can optionally comprise a nucleic acid sequence having complementarity to a sense region of the siNA molecule.

By "target nucleic acid" or "target polynucleotide" is meant any nucleic acid sequence whose expression or activity is to be modulated. The target nucleic acid can be DNA or RNA.

By "complementarity" is meant that a nucleic acid can form hydrogen bond(s) with another nucleic acid sequence by either traditional Watson-Crick or other non-traditional types. In reference to the nucleic molecules of the present invention, the binding free energy for a nucleic acid molecule with its complementary sequence is sufficient to allow the relevant function of the nucleic acid to proceed, e.g., RNAi activity. Determination of binding free energies for nucleic acid molecules is well known in the

art (see, e.g., Turner *et al.*, 1987, *CSH Symp. Quant. Biol.* LII pp.123-133; Frier *et al.*, 1986, *Proc. Nat. Acad. Sci. USA* 83:9373-9377; Turner *et al.*, 1987, *J. Am. Chem. Soc.* 109:3783-3785). A percent complementarity indicates the percentage of contiguous residues in a nucleic acid molecule that can form hydrogen bonds (e.g., Watson-Crick base pairing) with a second nucleic acid sequence (e.g., 5, 6, 7, 8, 9, or 10 nucleotides out of a total of 10 nucleotides in the first oligonucleotide being based paired to a second nucleic acid sequence having 10 nucleotides represents 50%, 60%, 70%, 80%, 90%, and 100% complementary respectively). "Perfectly complementary" means that all the contiguous residues of a nucleic acid sequence will hydrogen bond with the same number of contiguous residues in a second nucleic acid sequence. In one embodiment, a siNA molecule of the invention comprises about 15 to about 30 or more (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30 or more) nucleotides that are complementary to one or more target nucleic acid molecules or a portion thereof.

In one embodiment, siNA molecules of the invention that down regulate or reduce target gene expression are used for preventing or treating diseases, traits, disorders, and/or conditions in a subject or organism.

By "proliferative disease" or "cancer" as used herein is meant, any disease, condition, trait, genotype or phenotype characterized by unregulated cell growth or replication as is known in the art, including AIDS related cancers such as Kaposi's sarcoma; breast cancers; bone cancers such as Osteosarcoma, Chondrosarcomas, Ewing's sarcoma, Fibrosarcomas, Giant cell tumors, Adamantinomas, and Chordomas; Brain cancers such as Meningiomas, Glioblastomas, Lower-Grade Astrocytomas, Oligodendrocytomas, Pituitary Tumors, Schwannomas, and Metastatic brain cancers; cancers of the head and neck including various lymphomas such as mantle cell lymphoma, non-Hodgkins lymphoma, adenoma, squamous cell carcinoma, laryngeal carcinoma, gallbladder and bile duct cancers, cancers of the retina such as retinoblastoma, cancers of the esophagus, gastric cancers, multiple myeloma, ovarian cancer, uterine cancer, thyroid cancer, testicular cancer, endometrial cancer, melanoma, colorectal cancer, lung cancer, bladder cancer, prostate cancer, lung cancer (including non-small cell lung carcinoma), pancreatic cancer, sarcomas, Wilms' tumor, cervical cancer, head and neck cancer, skin cancers, nasopharyngeal carcinoma, liposarcoma,

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epithelial carcinoma, renal cell carcinoma, gallbladder adeno carcinoma, parotid adenocarcinoma, endometrial sarcoma, multidrug resistant cancers, and leukemias such as acute myelogenous leukemia (AML), chronic myelogenous leukemia (CML), acute lymphocytic leukemia (ALL), and chronic lymphocytic leukemia,; and proliferative diseases and conditions, such as neovascularization associated with tumor angiogenesis, macular degeneration (e.g., wet/dry AMD), corneal neovascularization, diabetic retinopathy, neovascular glaucoma, myopic degeneration and other proliferative diseases and conditions such as restenosis and polycystic kidney disease, and any other cancer or proliferative disease, condition, trait, genotype or phenotype that can respond to the modulation of disease related gene expression in a cell or tissue, alone or in combination with other therapies.

By "inflammatory disease" or "inflammatory condition" as used herein is meant any disease, condition, trait, genotype or phenotype characterized by an inflammatory or allergic process as is known in the art, such as inflammation, acute inflammation, chronic inflammation, respiratory disease, atherosclerosis, restenosis, asthma, allergic rhinitis, atopic dermatitis, septic shock, rheumatoid arthritis, inflammatory bowel disease, inflammatory pelvic disease, pain, ocular inflammatory disease, celiac disease, Leigh Syndrome, Glycerol Kinase Deficiency, Familial eosinophilia (FE), autosomal recessive spastic ataxia, laryngeal inflammatory disease; Tuberculosis, Chronic cholecystitis, Bronchiectasis, Silicosis and other pneumoconioses, and any other inflammatory disease, condition, trait, genotype or phenotype that can respond to the modulation of disease related gene expression in a cell or tissue, alone or in combination with other therapies.

By "autoimmune disease" or "autoimmune condition" as used herein is meant, any disease, condition, trait, genotype or phenotype characterized by autoimmunity as is known in the art, such as multiple sclerosis, diabetes mellitus, lupus, celiac disease, Crohn's disease, ulcerative colitis, Guillain-Barre syndrome, scleroderms, Goodpasture's syndrome, Wegener's granulomatosis, autoimmune epilepsy, Rasmussen's encephalitis, Primary biliary sclerosis, Sclerosing cholangitis, Autoimmune hepatitis, Addison's disease, Hashimoto's thyroiditis, Fibromyalgia, Menier's syndrome; transplantation rejection (e.g., prevention of allograft rejection) pernicious anemia, rheumatoid arthritis, systemic lupus erythematosus, dermatomyositis, Sjogren's syndrome, lupus

erythematous, multiple sclerosis, myasthenia gravis, Reiter's syndrome, Grave's disease, and any other autoimmune disease, condition, trait, genotype or phenotype that can respond to the modulation of disease related gene expression in a cell or tissue, alone or in combination with other therapies.

- 5 By "infectious disease" is meant any disease, condition, trait, genotype or phenotype associated with an infectious agent, such as a virus, bacteria, fungus, prion, or parasite. Non-limiting examples of various viral genes that can be targeted using siNA molecules of the invention include Hepatitis C Virus (HCV, for example Genbank Accession Nos: D11168, D50483.1, L38318 and S82227), Hepatitis B Virus (HBV, for
- 10 example GenBank Accession No. AF100308.1), Human Immunodeficiency Virus type 1 (HIV-1, for example GenBank Accession No. U51188), Human Immunodeficiency Virus type 2 (HIV-2, for example GenBank Accession No. X60667), West Nile Virus (WNV for example GenBank accession No. NC_001563), cytomegalovirus (CMV for example GenBank Accession No. NC_001347), respiratory syncytial virus (RSV for
- 15 example GenBank Accession No. NC_001781), influenza virus (for example GenBank Accession No. AF037412, rhinovirus (for example, GenBank accession numbers: D00239, X02316, X01087, L24917, M16248, K02121, X01087), papillomavirus (for example GenBank Accession No. NC_001353), Herpes Simplex Virus (HSV for example GenBank Accession No. NC_001345), and other viruses such as HTLV (for
- 20 example GenBank Accession No. AJ430458). Due to the high sequence variability of many viral genomes, selection of siNA molecules for broad therapeutic applications would likely involve the conserved regions of the viral genome. Nonlimiting examples of conserved regions of the viral genomes include but are not limited to 5'-Non Coding Regions (NCR), 3'- Non Coding Regions (NCR) and/or internal ribosome entry sites
- 25 (IRES). siNA molecules designed against conserved regions of various viral genomes will enable efficient inhibition of viral replication in diverse patient populations and may ensure the effectiveness of the siNA molecules against viral quasi species which evolve due to mutations in the non-conserved regions of the viral genome. Non-limiting examples of bacterial infections include Actinomycosis, Anthrax, Aspergillosis,
- 30 Bacteremia, Bacterial Infections and Mycoses, Bartonella Infections, Botulism, Brucellosis, Burkholderia Infections, Campylobacter Infections, Candidiasis, Cat-Scratch Disease, Chlamydia Infections, Cholera , Clostridium Infections, Coccidioidomycosis,

Cross Infection, Cryptococcosis, Dermatomyces, Dermatomyces, Diphtheria, Ehrlichiosis, Escherichia coli Infections, Fasciitis, Necrotizing, Fusobacterium Infections, Gas Gangrene, Gram-Negative Bacterial Infections, Gram-Positive Bacterial Infections, Histoplasmosis, Impetigo, Klebsiella Infections, Legionellosis, Leprosy, 5 Leptospirosis, Listeria Infections, Lyme Disease, Maduromycosis, Melioidosis, Mycobacterium Infections, Mycoplasma Infections, Mycoses, Nocardia Infections, Onychomycosis, Ornithosis, Plague, Pneumococcal Infections, Pseudomonas Infections, Q Fever, Rat-Bite Fever, Relapsing Fever, Rheumatic Fever, Rickettsia Infections, Rocky Mountain Spotted Fever, Salmonella Infections, Scarlet Fever, Scrub Typhus, 10 Sepsis, Sexually Transmitted Diseases - Bacterial, Bacterial Skin Diseases, Staphylococcal Infections, Streptococcal Infections, Tetanus, Tick-Borne Diseases, Tuberculosis, Tularemia, Typhoid Fever, Typhus, Epidemic Louse-Borne, Vibrio Infections, Yaws, Yersinia Infections, Zoonoses, and Zygomycosis. Non-limiting examples of fungal infections include Aspergillosis, Blastomycosis, 15 Coccidioidomycosis, Cryptococcosis, Fungal Infections of Fingernails and Toenails, Fungal Sinusitis, Histoplasmosis, Histoplasmosis, Mucormycosis, Nail Fungal Infection, Paracoccidioidomycosis, Sporotrichosis, Valley Fever (Coccidioidomycosis), and Mold Allergy.

By "neurologic disease" or "neurological disease" is meant any disease, disorder, 20 or condition affecting the central or peripheral nervous system, including ADHD, AIDS - Neurological Complications, Absence of the Septum Pellucidum, Acquired Epileptiform Aphasia, Acute Disseminated Encephalomyelitis, Adrenoleukodystrophy, Agenesis of the Corpus Callosum, Agnosia, Aicardi Syndrome, Alexander Disease, Alpers' Disease, Alternating Hemiplegia, Alzheimer's Disease, Amyotrophic Lateral Sclerosis, 25 Anencephaly, Aneurysm, Angelman Syndrome, Angiomatosis, Anoxia, Aphasia, Apraxia, Arachnoid Cysts, Arachnoiditis, Arnold-Chiari Malformation, Arteriovenous Malformation, Aspartame, Asperger Syndrome, Ataxia Telangiectasia, Ataxia, Attention Deficit-Hyperactivity Disorder, Autism, Autonomic Dysfunction, Back Pain, Barth Syndrome, Batten Disease, Behcet's Disease, Bell's Palsy, Benign Essential 30 Blepharospasm, Benign Focal Amyotrophy, Benign Intracranial Hypertension, Bernhardt-Roth Syndrome, Binswanger's Disease, Blepharospasm, Bloch-Sulzberger Syndrome, Brachial Plexus Birth Injuries, Brachial Plexus Injuries, Bradbury-Eggleston

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- Syndrome, Brain Aneurysm, Brain Injury, Brain and Spinal Tumors, Brown-Sequard Syndrome, Bulbospinal Muscular Atrophy, Canavan Disease, Carpal Tunnel Syndrome, Causalgia, Cavernomas, Cavernous Angioma, Cavernous Malformation, Central Cervical Cord Syndrome, Central Cord Syndrome, Central Pain Syndrome, Cephalic
- 5 Disorders, Cerebellar Degeneration, Cerebellar Hypoplasia, Cerebral Aneurysm, Cerebral Arteriosclerosis, Cerebral Atrophy, Cerebral Beriberi, Cerebral Gigantism, Cerebral Hypoxia, Cerebral Palsy, Cerebro-Oculo-Facio-Skeletal Syndrome, Charcot-Marie-Tooth Disorder, Chiari Malformation, Chorea, Choreoacanthocytosis, Chronic Inflammatory Demyelinating Polyneuropathy (CIDP), Chronic Orthostatic Intolerance,
- 10 Chronic Pain, Cockayne Syndrome Type II, Coffin Lowry Syndrome, Coma, including Persistent Vegetative State, Complex Regional Pain Syndrome, Congenital Facial Diplegia, Congenital Myasthenia, Congenital Myopathy, Congenital Vascular Cavernous Malformations, Corticobasal Degeneration, Cranial Arteritis, Craniosynostosis, Creutzfeldt-Jakob Disease, Cumulative Trauma Disorders, Cushing's Syndrome,
- 15 Cytomegalic Inclusion Body Disease (CIBD), Cytomegalovirus Infection, Dancing Eyes-Dancing Feet Syndrome, Dandy-Walker Syndrome, Dawson Disease, De Morsier's Syndrome, Dejerine-Klumpke Palsy, Dementia - Multi-Infarct, Dementia - Subcortical, Dementia With Lewy Bodies, Dermatomyositis, Developmental Dyspraxia, Devic's Syndrome, Diabetic Neuropathy, Diffuse Sclerosis, Dravet's Syndrome, Dysautonomia,
- 20 Dysgraphia, Dyslexia, Dysphagia, Dyspraxia, Dystonias, Early Infantile Epileptic Encephalopathy, Empty Sella Syndrome, Encephalitis Lethargica, Encephalitis and Meningitis, Encephaloceles, Encephalopathy, Encephalotrigeminal Angiomatosis, Epilepsy, Erb's Palsy, Erb-Duchenne and Dejerine-Klumpke Palsies, Fabry's Disease, Fahr's Syndrome, Fainting, Familial Dysautonomia, Familial Hemangioma, Familial
- 25 Idiopathic Basal Ganglia Calcification, Familial Spastic Paralysis, Febrile Seizures (e.g., GEFS and GEFS plus), Fisher Syndrome, Floppy Infant Syndrome, Friedrich's Ataxia, Gaucher's Disease, Gerstmann's Syndrome, Gerstmann-Straussler-Scheinker Disease, Giant Cell Arteritis, Giant Cell Inclusion Disease, Globoid Cell Leukodystrophy, Glossopharyngeal Neuralgia, Guillain-Barre Syndrome, HTLV-1 Associated
- 30 Myelopathy, Hallervorden-Spatz Disease, Head Injury, Headache, Hemispheric Continuity, Hemifacial Spasm, Hemiplegia Alterans, Hereditary Neuropathies, Hereditary Spastic Paraplegia, Hereditary Ataxia Polyneuriticiformis, Herpes Zoster Oticus, Herpes

- Zoster, Hirayama Syndrome, Holoprosencephaly, Huntington's Disease, Hydranencephaly, Hydrocephalus - Normal Pressure, Hydrocephalus, Hydromyelia, Hypercortisolism, Hypersomnia, Hypertonia, Hypotonia, Hypoxia, Immune-Mediated Encephalomyelitis, Inclusion Body Myositis, Incontinentia Pigmenti, Infantile
- 5 Hypotonia, Infantile Phytanic Acid Storage Disease, Infantile Refsum Disease, Infantile Spasms, Inflammatory Myopathy, Intestinal Lipodystrophy, Intracranial Cysts, Intracranial Hypertension, Isaac's Syndrome, Joubert Syndrome, Kearns-Sayre Syndrome, Kennedy's Disease, Kinsbourne syndrome, Kleine-Levin syndrome, Klippel Feil Syndrome, Klippel-Trenaunay Syndrome (KTS), Klüver-Bucy Syndrome,
- 10 Korsakoff's Amnesic Syndrome, Krabbe Disease, Kugelberg-Welander Disease, Kuru, Lambert-Eaton Myasthenic Syndrome, Landau-Kleffner Syndrome, Lateral Femoral Cutaneous Nerve Entrapment, Lateral Medullary Syndrome, Learning Disabilities, Leigh's Disease, Lennox-Gastaut Syndrome, Lesch-Nyhan Syndrome, Leukodystrophy, Levine-Critchley Syndrome, Lewy Body Dementia, Lissencephaly, Locked-In
- 15 Syndrome, Lou Gehrig's Disease, Lupus - Neurological Sequelae, Lyme Disease - Neurological Complications, Machado-Joseph Disease, Macrencephaly, Megalencephaly, Melkersson-Rosenthal Syndrome, Meningitis, Menkes Disease, Meralgia Paresthetica, Metachromatic Leukodystrophy, Microcephaly, Migraine, Miller Fisher Syndrome, Mini-Stroke, Mitochondrial Myopathies, Mobius Syndrome,
- 20 Monomelic Amyotrophy, Motor Neuron Diseases, Moyamoya Disease, Mucopolidoses, Mucopolysaccharidoses, Multi-Infarct Dementia, Multifocal Motor Neuropathy, Multiple Sclerosis, Multiple System Atrophy with Orthostatic Hypotension, Multiple System Atrophy, Muscular Dystrophy, Myasthenia - Congenital, Myasthenia Gravis, Myelinoclastic Diffuse Sclerosis, Myoclonic Encephalopathy of Infants, Myoclonus,
- 25 Myopathy - Congenital, Myopathy - Thyrotoxic, Myopathy, Myotonia Congenita, Myotonia, Narcolepsy, Neuroacanthocytosis, Neurodegeneration with Brain Iron Accumulation, Neurofibromatosis, Neuroleptic Malignant Syndrome, Neurological Complications of AIDS, Neurological Manifestations of Pompe Disease, Neuromyelitis Optica, Neuromyotonia, Neuronal Ceroid Lipofuscinosis, Neuronal Migration Disorders,
- 30 Neuropathy - Hereditary, Neurosarcooidosis, Neurotoxicity, Nevus Cavemosus, Niemann-Pick Disease, O'Sullivan-McLeod Syndrome, Occipital Neuralgia, Occult Spinal Dysraphism Sequence, Ohtahara Syndrome, Olivopontocerebellar Atrophy, Opsoclonus

- Myoclonus, Orthostatic Hypotension, Overuse Syndrome, Pain - Chronic, Paraneoplastic Syndromes, Paresthesia, Parkinson's Disease, Parmyotonia Congenita, Paroxysmal Choreoathetosis, Paroxysmal Hemicrania, Parry-Romberg, Pelizaeus-Merzbacher Disease, Pena Shokeir II Syndrome, Perineural Cysts, Periodic Paralysis, Peripheral
- 5 Neuropathy, Periventricular Leukomalacia, Persistent Vegetative State, Pervasive Developmental Disorders, Phytanic Acid Storage Disease, Pick's Disease, Piriformis Syndrome, Pituitary Tumors, Polymyositis, Pompe Disease, Porencephaly, Post-Polio Syndrome, Postherpetic Neuralgia, Postinfectious Encephalomyelitis, Postural Hypotension, Postural Orthostatic Tachycardia Syndrome, Postural Tachycardia
- 10 Syndrome, Primary Lateral Sclerosis, Prion Diseases, Progressive Hemifacial Atrophy, Progressive Locomotor Ataxia, Progressive Multifocal Leukoencephalopathy, Progressive Sclerosing Poliodystrophy, Progressive Supranuclear Palsy, Pseudotumor Cerebri, Pyridoxine Dependent and Pyridoxine Responsive Seizure Disorders, Ramsay Hunt Syndrome Type I, Ramsay Hunt Syndrome Type II, Rasmussen's Encephalitis and
- 15 other autoimmune epilepsies, Reflex Sympathetic Dystrophy Syndrome, Refsum Disease - Infantile, Refsum Disease, Repetitive Motion Disorders, Repetitive Stress Injuries, Restless Legs Syndrome, Retrovirus-Associated Myelopathy, Rett Syndrome, Reye's Syndrome, Riley-Day Syndrome, SUNCT Headache, Sacral Nerve Root Cysts, Saint Vitus Dance, Salivary Gland Disease, Sandhoff Disease, Schilder's Disease,
- 20 Schizencephaly, Seizure Disorders, Septo-Optic Dysplasia, Severe Myoclonic Epilepsy of Infancy (SMEI), Shaken Baby Syndrome, Shingles, Shy-Drager Syndrome, Sjogren's Syndrome, Sleep Apnea, Sleeping Sickness, Soto's Syndrome, Spasticity, Spina Bifida, Spinal Cord Infarction, Spinal Cord Injury, Spinal Cord Tumors, Spinal Muscular Atrophy, Spinocerebellar Atrophy, Steele-Richardson-Olszewski Syndrome, Stiff-Person
- 25 Syndrome, Striatonigral Degeneration, Stroke, Sturge-Weber Syndrome, Subacute Sclerosing Panencephalitis, Subcortical Arteriosclerotic Encephalopathy, Swallowing Disorders, Sydenham Chorea, Syncope, Syphilitic Spinal Sclerosis, Syringohydromyelia, Syringomyelia, Systemic Lupus Erythematosus, Tabes Dorsalis, Tardive Dyskinesia, Tarlov Cysts, Tay-Sachs Disease, Temporal Arteritis, Tethered Spinal Cord Syndrome,
- 30 Thomsen Disease, Thoracic Outlet Syndrome, Thyrotoxic Myopathy, Tic Douloureux, Todd's Paralysis, Tourette Syndrome, Transient Ischemic Attack, Transmissible Spongiform Encephalopathies, Transverse Myelitis, Traumatic Brain Injury, Tremor,

- Trigeminal Neuralgia, Tropical Spastic Paraparesis, Tuberous Sclerosis, Vascular Erectile Tumor, Vasculitis including Temporal Arteritis, Von Economo's Disease, Von Hippel-Lindau disease (VHL), Von Recklinghausen's Disease, Wallenberg's Syndrome, Werdnig-Hoffman Disease, Wernicke-Korsakoff Syndrome, West Syndrome, Whipple's Disease, Williams Syndrome, Wilson's Disease, X-Linked Spinal and Bulbar Muscular Atrophy, and Zellweger Syndrome.

By "respiratory disease" is meant, any disease or condition affecting the respiratory tract, such as asthma, chronic obstructive pulmonary disease or "COPD", allergic rhinitis, sinusitis, pulmonary vasoconstriction, inflammation, allergies, impeded respiration, respiratory distress syndrome, cystic fibrosis, pulmonary hypertension, pulmonary vasoconstriction, emphysema, and any other respiratory disease, condition, trait, genotype or phenotype that can respond to the modulation of disease related gene expression in a cell or tissue, alone or in combination with other therapies.

By "cardiovascular disease" is meant and disease or condition affecting the heart and vasculature, including but not limited to, coronary heart disease (CHD), cerebrovascular disease (CVD), aortic stenosis, peripheral vascular disease, atherosclerosis, arteriosclerosis, myocardial infarction (heart attack), cerebrovascular diseases (stroke), transient ischaemic attacks (TIA), angina (stable and unstable), atrial fibrillation, arrhythmia, valvular disease, congestive heart failure, hypercholesterolemia, type I hyperlipoproteinemia, type II hyperlipoproteinemia, type III hyperlipoproteinemia, type IV hyperlipoproteinemia, type V hyperlipoproteinemia, secondary hypertriglyceridemia, and familial lecithin cholesterol acyltransferase deficiency.

By "ocular disease" as used herein is meant, any disease, condition, trait, genotype or phenotype of the eye and related structures as is known in the art, such as Cystoid Macular Edema, Asteroid Hyalosis, Pathological Myopia and Posterior Staphyloma, Toxocariasis (Ocular Larva Migrans), Retinal Vein Occlusion, Posterior Vitreous Detachment, Tractional Retinal Tears, Epiretinal Membrane, Diabetic Retinopathy, Lattice Degeneration, Retinal Vein Occlusion, Retinal Artery Occlusion, Macular Degeneration (e.g., age related macular degeneration such as wet AMD or dry AMD), Toxoplasmosis, Choroidal Melanoma, Acquired Retinoschisis, Hollenhorst Plaque, Idiopathic Central Serous Chorioretinopathy, Macular Hole, Presumed Ocular

- Histoplasmosis Syndrome, Retinal Macroaneurysm, Retinitis Pigmentosa, Retinal Detachment, Hypertensive Retinopathy, Retinal Pigment Epithelium (RPE) Detachment, Papillophlebitis, Ocular Ischemic Syndrome, Coats' Disease, Leber's Miliary Aneurysm, Conjunctival Neoplasms, Allergic Conjunctivitis, Vernal Conjunctivitis, Acute Bacterial
- 5 Conjunctivitis, Allergic Conjunctivitis & Vernal Keratoconjunctivitis, Viral Conjunctivitis, Bacterial Conjunctivitis, Chlamydial & Gonococcal Conjunctivitis, Conjunctival Laceration, Episcleritis, Scleritis, Pingueculitis, Pterygium, Superior Limbic Keratoconjunctivitis (SLK of Theodore), Toxic Conjunctivitis, Conjunctivitis with Pseudomembrane, Giant Papillary Conjunctivitis, Terrien's Marginal Degeneration,
- 10 Acanthamoeba Keratitis, Fungal Keratitis, Filamentary Keratitis, Bacterial Keratitis, Keratitis Sicca/Dry Eye Syndrome, Bacterial Keratitis, Herpes Simplex Keratitis, Sterile Corneal Infiltrates, Phlyctenulosis, Corneal Abrasion & Recurrent Corneal Erosion, Corneal Foreign Body, Chemical Burs, Epithelial Basement Membrane Dystrophy (EBMD), Thygeson's Superficial Punctate Keratopathy, Corneal Laceration, Salzmann's
- 15 Nodular Degeneration, Fuchs' Endothelial Dystrophy, Crystalline Lens Subluxation, Ciliary-Block Glaucoma, Primary Open-Angle Glaucoma, Pigment Dispersion Syndrome and Pigmentary Glaucoma, Pseudoexfoliation Syndrom and Pseudoexfoliative Glaucoma, Anterior Uveitis, Primary Open Angle Glaucoma, Uveitic Glaucoma & Glaucomatocyclitic Crisis, Pigment Dispersion Syndrome & Pigmentary Glaucoma,
- 20 Acute Angle Closure Glaucoma, Anterior Uveitis, Hyphema, Angle Recession Glaucoma, Lens Induced Glaucoma, Pseudoexfoliation Syndrome and Pseudoexfoliative Glaucoma, Axenfeld-Rieger Syndrome, Neovascular Glaucoma, Pars Planitis, Choroidal Rupture, Duane's Retraction Syndrome, Toxic/Nutritional Optic Neuropathy, Aberrant Regeneration of Cranial Nerve III, Intracranial Mass Lesions, Carotid-Cavernous Sinus
- 25 Fistula, Anterior Ischemic Optic Neuropathy, Optic Disc Edema & Papilledema, Cranial Nerve III Palsy, Cranial Nerve IV Palsy, Cranial Nerve VI Palsy, Cranial Nerve VII (Facial Nerve) Palsy, Horner's Syndrome, Internuclear Ophthalmoplegia, Optic Nerve Head Hypoplasia, Optic Pit, Tonic Pupil, Optic Nerve Head Drusen, Demyelinating Optic Neuropathy (Optic Neuritis, Retrobulbar Optic Neuritis), Amaurosis Fugax and
- 30 Transient Ischemic Attack, Pseudotumor Cerebri, Pituitary Adenoma, Molluscum Contagiosum, Canaliculitis, Verruca and Papilloma, Pediculosis and Phthiriasis, Blepharitis, Hordeolum, Preseptal Cellulitis, Chalazion, Basal Cell Carcinoma, Herpes

Zoster Ophthalmicus, Pediculosis & Phthiriasis, Blow-out Fracture, Chronic Epiphora, Dacryocystitis, Herpes Simplex Blepharitis, Orbital Cellulitis, Senile Entropion, and Squamous Cell Carcinoma.

By "metabolic disease" is meant any disease or condition affecting metabolic pathways as in known in the art. Metabolic disease can result in an abnormal metabolic process, either congenital due to inherited enzyme abnormality (inborn errors of metabolism) or acquired due to disease of an endocrine organ or failure of a metabolically important organ such as the liver. In one embodiment, metabolic disease includes obesity, insulin resistance, and diabetes (e.g., type I and/or type II diabetes).

In one embodiment of the present invention, each sequence of a siRNA molecule of the invention is independently about 15 to about 30 nucleotides in length, in specific embodiments about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30 nucleotides in length. In another embodiment, the siRNA duplexes of the invention independently comprise about 15 to about 30 base pairs (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30). In another embodiment, one or more strands of the siRNA molecule of the invention independently comprises about 15 to about 30 nucleotides (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30) that are complementary to a target nucleic acid molecule. In yet another embodiment, siRNA molecules of the invention comprising hairpin or circular structures are about 35 to about 55 (e.g., about 35, 40, 45, 50 or 55) nucleotides in length, or about 38 to about 44 (e.g., about 38, 39, 40, 41, 42, 43, or 44) nucleotides in length and comprising about 15 to about 25 (e.g., about 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, or 25) base pairs. Exemplary siRNA molecules of the invention are shown in Figures 4-5.

As used herein "cell" is used in its usual biological sense, and does not refer to an entire multicellular organism, e.g., specifically does not refer to a human. The cell can be present in an organism, e.g., birds, plants and mammals such as humans, cows, sheep, apes, monkeys, swine, dogs, and cats. The cell can be prokaryotic (e.g., bacterial cell) or eukaryotic (e.g., mammalian or plant cell). The cell can be of somatic or germ line origin, totipotent or pluripotent, dividing or non-dividing. The cell can also be derived from or can comprise a gamete or embryo, a stem cell, or a fully differentiated cell.

The siNA molecules of the invention are added directly, or can be complexed with cationic lipids, packaged within liposomes, or otherwise delivered to target cells or tissues. The nucleic acid or nucleic acid complexes can be locally administered to relevant tissues *ex vivo*, or *in vivo* through direct dermal application, transdermal application, or injection, with or without their incorporation in biopolymers.

In another aspect, the invention provides mammalian cells containing one or more siNA molecules of this invention. The one or more siNA molecules can independently be targeted to the same or different sites.

By "RNA" is meant a molecule comprising at least one ribonucleotide residue. By "ribonucleotide" is meant a nucleotide with a hydroxyl group at the 2' position of a β -D-ribofuranose moiety. The terms include double-stranded RNA, single-stranded RNA, isolated RNA such as partially purified RNA, essentially pure RNA, synthetic RNA, recombinantly produced RNA, as well as altered RNA that differs from naturally occurring RNA by the addition, deletion, substitution and/or alteration of one or more nucleotides. Such alterations can include addition of non-nucleotide material, such as to the end(s) of the siNA or internally, for example at one or more nucleotides of the RNA. Nucleotides in the RNA molecules of the instant invention can also comprise non-standard nucleotides, such as non-naturally occurring nucleotides or chemically synthesized nucleotides or deoxynucleotides. These altered RNAs can be referred to as analogs or analogs of naturally-occurring RNA.

By "subject" is meant an organism, which is a donor or recipient of explanted cells or the cells themselves. "Subject" also refers to an organism to which the nucleic acid molecules of the invention can be administered. A subject can be a mammal or mammalian cells, including a human or human cells.

The term "phosphorothioate" as used herein refers to an internucleotide linkage having Formula I, wherein Z and/or W comprise a sulfur atom. Hence, the term phosphorothioate refers to both phosphorothioate and phosphorodithioate internucleotide linkages.

The term "phosphonoacetate" as used herein refers to an internucleotide linkage having Formula I, wherein Z and/or W comprise an acetyl or protected acetyl group.

The term "thiophosphonoacetate" as used herein refers to an internucleotide linkage having Formula I, wherein Z comprises an acetyl or protected acetyl group and
5 W comprises a sulfur atom or alternately W comprises an acetyl or protected acetyl group and Z comprises a sulfur atom.

The term "universal base" as used herein refers to nucleotide base analogs that form base pairs with each of the natural DNA/RNA bases with little discrimination between them. Non-limiting examples of universal bases include C-phenyl, C-naphthyl
10 and other aromatic derivatives, inosine, azole carboxamides, and nitroazole derivatives such as 3-nitropyrrole, 4-nitroindole, 5-nitroindole, and 6-nitroindole as known in the art (see for example Loakes, 2001, *Nucleic Acids Research*, 29, 2437-2447).

The term "acyclic nucleotide" as used herein refers to any nucleotide having an acyclic ribose sugar, for example where any of the ribose carbons (C1, C2, C3, C4, or
15 C5), are independently or in combination absent from the nucleotide.

The nucleic acid molecules of the instant invention, individually, or in combination or in conjunction with other drugs, can be used to for preventing or treating diseases, traits, disorders, and/or conditions described herein or otherwise known in the art, in a subject or organism. For example, the siNA molecules can be administered to a subject
20 or can be administered to other appropriate cells evident to those skilled in the art, individually or in combination with one or more drugs under conditions suitable for the treatment.

In a further embodiment, the siNA molecules can be used in combination with other known treatments to prevent or treat preventing or treating diseases, traits,
25 disorders, and/or conditions described herein or otherwise known in the art, in a subject or organism. For example, the described molecules could be used in combination with one or more known compounds, treatments, or procedures to prevent or treat diseases, traits, disorders, and/or conditions described herein or otherwise known in the art, in a subject or organism.

In one embodiment, the invention features an expression vector comprising a nucleic acid sequence encoding at least one siNA molecule of the invention, in a manner which allows expression of the siNA molecule. For example, the vector can contain sequence(s) encoding both strands of a siNA molecule comprising a duplex. The vector
5 can also contain sequence(s) encoding a single nucleic acid molecule that is self-complementary and thus forms a siNA molecule. Non-limiting examples of such expression vectors are described in Paul *et al.*, 2002, *Nature Biotechnology*, 19, 505; Miyagishi and Taira, 2002, *Nature Biotechnology*, 19, 497; Lee *et al.*, 2002, *Nature Biotechnology*, 19, 500; and Novina *et al.*, 2002, *Nature Medicine*, advance online
10 publication doi:10.1038/nm725.

In another embodiment, the invention features a mammalian cell, for example, a human cell, including an expression vector of the invention.

In yet another embodiment, the expression vector of the invention comprises a sequence for a siNA molecule having complementarity to a RNA molecule referred to by
15 Genbank Accession numbers, for example Genbank Accession Nos. shown in Table I.

In one embodiment, an expression vector of the invention comprises a nucleic acid sequence encoding two or more siNA molecules, which can be the same or different.

In another aspect of the invention, siNA molecules that interact with target RNA molecules and down-regulate gene encoding target RNA molecules (for example target
20 RNA molecules referred to by Genbank Accession numbers herein) are expressed from transcription units inserted into DNA or RNA vectors. The recombinant vectors can be DNA plasmids or viral vectors. siNA expressing viral vectors can be constructed based on, but not limited to, adeno-associated virus, retrovirus, adenovirus, or alphavirus. The recombinant vectors capable of expressing the siNA molecules can be delivered as
25 described herein, and persist in target cells. Alternatively, viral vectors can be used that provide for transient expression of siNA molecules. Such vectors can be repeatedly administered as necessary. Once expressed, the siNA molecules bind and down-regulate gene function or expression via RNA interference (RNAi). Delivery of siNA expressing vectors can be systemic, such as by intravenous or intramuscular administration, by
30 administration to target cells ex-planted from a subject followed by reintroduction into

the subject, or by any other means that would allow for introduction into the desired target cell.

By "vectors" is meant any nucleic acid- and/or viral-based technique used to deliver a desired nucleic acid.

- 5 Other features and advantages of the invention will be apparent from the following description of the preferred embodiments thereof, and from the claims.

BRIEF DESCRIPTION OF THE DRAWINGS

- Figure 1 shows a non-limiting example of a scheme for the synthesis of siNA molecules. The complementary siNA sequence strands, strand 1 and strand 2, are synthesized in tandem and are connected by a cleavable linkage, such as a nucleotide succinate or abasic succinate, which can be the same or different from the cleavable linker used for solid phase synthesis on a solid support. The synthesis can be either solid phase or solution phase, in the example shown, the synthesis is a solid phase synthesis. The synthesis is performed such that a protecting group, such as a dimethoxytrityl group, remains intact on the terminal nucleotide of the tandem oligonucleotide. Upon cleavage and deprotection of the oligonucleotide, the two siNA strands spontaneously hybridize to form a siNA duplex, which allows the purification of the duplex by utilizing the properties of the terminal protecting group, for example by applying a trityl on purification method wherein only duplexes/oligonucleotides with the terminal protecting group are isolated.
- 10
15
20

- Figure 2 shows a MALDI-TOF mass spectrum of a purified siNA duplex synthesized by a method of the invention. The two peaks shown correspond to the predicted mass of the separate siNA sequence strands. This result demonstrates that the siNA duplex generated from tandem synthesis can be purified as a single entity using a simple trityl-on purification methodology.
- 25

Figure 3 shows a non-limiting proposed mechanistic representation of target RNA degradation involved in RNAi. Double-stranded RNA (dsRNA), which is generated by RNA-dependent RNA polymerase (RdRP) from foreign single-stranded RNA, for example viral, transposon, or other exogenous RNA, activates the DICER enzyme that in

turn generates siRNA duplexes. Alternately, synthetic or expressed siRNA can be introduced directly into a cell by appropriate means. An active siRNA complex forms which recognizes a target RNA, resulting in degradation of the target RNA by the RISC endonuclease complex or in the synthesis of additional RNA by RNA-dependent RNA polymerase (RdRP), which can activate DICER and result in additional siRNA molecules, thereby amplifying the RNAi response.

Figure 4A-F shows non-limiting examples of chemically-modified siRNA constructs of the present invention. In the figure, N stands for any nucleotide (adenosine, guanosine, cytosine, uridine, or optionally thymidine, for example thymidine can be substituted in the overhanging regions designated by parenthesis (N N). Various modifications are shown for the sense and antisense strands of the siRNA constructs.

Figure 4A: The sense strand comprises 21 nucleotides wherein the two terminal 3'-nucleotides are optionally base paired and wherein all nucleotides present are ribonucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and wherein all nucleotides present are ribonucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. A modified internucleotide linkage, such as a phosphorothioate, phosphorodithioate or other modified internucleotide linkage as described herein, shown as "s", optionally connects the (N N) nucleotides in the antisense strand.

Figure 4B: The sense strand comprises 21 nucleotides wherein the two terminal 3'-nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides and all purine nucleotides that may be present are 2'-O-methyl modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and wherein all

pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides and all purine nucleotides that may be present are 2'-O-methyl modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. A modified internucleotide linkage, such as a phosphorothioate, phosphorodithioate or other modified internucleotide linkage as described herein, shown as "s", optionally connects the (N N) nucleotides in the sense and antisense strand.

Figure 4C: The sense strand comprises 21 nucleotides having 5'- and 3'- terminal cap moieties wherein the two terminal 3'-nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-O-methyl or 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. A modified internucleotide linkage, such as a phosphorothioate, phosphorodithioate or other modified internucleotide linkage as described herein, shown as "s", optionally connects the (N N) nucleotides in the antisense strand.

Figure 4D: The sense strand comprises 21 nucleotides having 5'- and 3'- terminal cap moieties wherein the two terminal 3'-nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein and wherein and all purine nucleotides that may be present are 2'-deoxy nucleotides. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides and all purine nucleotides that may be present are

2'-O-methyl modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. A modified internucleotide linkage, such as a phosphorothioate, phosphorodithioate or other modified internucleotide linkage as described herein, shown as "s", optionally connects the (N N) nucleotides in the antisense strand.

Figure 4E: The sense strand comprises 21 nucleotides having 5'- and 3'- terminal cap moieties wherein the two terminal 3'-nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides and all purine nucleotides that may be present are 2'-O-methyl modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. A modified internucleotide linkage, such as a phosphorothioate, phosphorodithioate or other modified internucleotide linkage as described herein, shown as "s", optionally connects the (N N) nucleotides in the antisense strand.

Figure 4F: The sense strand comprises 21 nucleotides having 5'- and 3'- terminal cap moieties wherein the two terminal 3'-nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein and wherein all purine nucleotides that may be present are 2'-deoxy nucleotides. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and having one 3'-terminal phosphorothioate internucleotide linkage and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides and all purine nucleotides that may be present are 2'-deoxy nucleotides except for (N N) nucleotides, which can comprise ribonucleotides,

deoxynucleotides, universal bases, or other chemical modifications described herein. A modified internucleotide linkage, such as a phosphorothioate, phosphorodithioate or other modified internucleotide linkage as described herein, shown as "s", optionally connects the (N N) nucleotides in the antisense strand. The antisense strand of constructs A-F comprise sequence complementary to any target nucleic acid sequence of the invention. Furthermore, when a glyceryl moiety (L) is present at the 3'-end of the antisense strand for any construct shown in Figure 4 A-F, the modified internucleotide linkage is optional.

Figure 5A-F shows non-limiting examples of specific chemically-modified siRNA sequences of the invention. A-F applies the chemical modifications described in Figure 4A-F to a target siRNA sequence. Such chemical modifications can be applied to any target sequence and/or target polymorphism sequence.

Figure 6 shows non-limiting examples of different siRNA constructs of the invention. The examples shown (constructs 1, 2, and 3) have 19 representative base pairs; however, different embodiments of the invention include any number of base pairs described herein. Bracketed regions represent nucleotide overhangs, for example, comprising about 1, 2, 3, or 4 nucleotides in length, preferably about 2 nucleotides. Constructs 1 and 2 can be used independently for RNAi activity. Construct 2 can comprise a polynucleotide or non-nucleotide linker, which can optionally be designed as a biodegradable linker. In one embodiment, the loop structure shown in construct 2 can comprise a biodegradable linker that results in the formation of construct 1 *in vivo* and/or *in vitro*. In another example, construct 3 can be used to generate construct 2 under the same principle wherein a linker is used to generate the active siRNA construct 2 *in vivo* and/or *in vitro*, which can optionally utilize another biodegradable linker to generate the active siRNA construct 1 *in vivo* and/or *in vitro*. As such, the stability and/or activity of the siRNA constructs can be modulated based on the design of the siRNA construct for use *in vivo* or *in vitro* and/or *in vitro*.

Figure 7A-C is a diagrammatic representation of a scheme utilized in generating an expression cassette to generate siRNA hairpin constructs.

Figure 7A: A DNA oligomer is synthesized with a 5'-restriction site (R1) sequence followed by a region having sequence identical (sense region of siNA) to a predetermined target sequence, wherein the sense region comprises, for example, about 19, 20, 21, or 22 nucleotides (N) in length, which is followed by a loop sequence of defined sequence (X), comprising, for example, about 3 to about 10 nucleotides.

Figure 7B: The synthetic construct is then extended by DNA polymerase to generate a hairpin structure having self-complementary sequence that will result in a siNA transcript having specificity for a target sequence and having self-complementary sense and antisense regions.

Figure 7C: The construct is heated (for example to about 95°C) to linearize the sequence, thus allowing extension of a complementary second DNA strand using a primer to the 3'-restriction sequence of the first strand. The double-stranded DNA is then inserted into an appropriate vector for expression in cells. The construct can be designed such that a 3'-terminal nucleotide overhang results from the transcription, for example, by engineering restriction sites and/or utilizing a poly-U termination region as described in Paul *et al.*, 2002, *Nature Biotechnology*, 29, 505-508.

Figure 8A-C is a diagrammatic representation of a scheme utilized in generating an expression cassette to generate double-stranded siNA constructs.

Figure 8A: A DNA oligomer is synthesized with a 5'-restriction (R1) site sequence followed by a region having sequence identical (sense region of siNA) to a predetermined target sequence, wherein the sense region comprises, for example, about 19, 20, 21, or 22 nucleotides (N) in length, and which is followed by a 3'-restriction site (R2) which is adjacent to a loop sequence of defined sequence (X).

Figure 8B: The synthetic construct is then extended by DNA polymerase to generate a hairpin structure having self-complementary sequence.

Figure 8C: The construct is processed by restriction enzymes specific to R1 and R2 to generate a double-stranded DNA which is then inserted into an appropriate vector for expression in cells. The transcription cassette is designed such that a U6 promoter region flanks each side of the dsDNA which generates the separate sense and antisense

strands of the siNA. Poly T termination sequences can be added to the constructs to generate U overhangs in the resulting transcript.

Figure 9A-E is a diagrammatic representation of a method used to determine target sites for siNA mediated RNAi within a particular target nucleic acid sequence, such as messenger RNA.

Figure 9A: A pool of siNA oligonucleotides are synthesized wherein the antisense region of the siNA constructs has complementarity to target sites across the target nucleic acid sequence, and wherein the sense region comprises sequence complementary to the antisense region of the siNA.

Figure 9B&C: (Figure 9B) The sequences are pooled and are inserted into vectors such that (Figure 9C) transfection of a vector into cells results in the expression of the siNA.

Figure 9D: Cells are sorted based on phenotypic change that is associated with modulation of the target nucleic acid sequence.

Figure 9E: The siNA is isolated from the sorted cells and is sequenced to identify efficacious target sites within the target nucleic acid sequence.

Figure 10 shows non-limiting examples of different stabilization chemistries (1-10) that can be used, for example, to stabilize the 3'-end of siNA sequences of the invention, including (1) [3'-3']-inverted deoxyribose; (2) deoxyribonucleotide; (3) [5'-3']-3'-deoxyribonucleotide; (4) [5'-3']-ribonucleotide; (5) [5'-3']-3'-O-methyl ribonucleotide; (6) 3'-glyceryl; (7) [3'-5']-3'-deoxyribonucleotide; (8) [3'-3']-deoxyribonucleotide; (9) [5'-2']-deoxyribonucleotide; and (10) [5'-3']-dideoxyribonucleotide. In addition to modified and unmodified backbone chemistries indicated in the figure, these chemistries can be combined with different backbone modifications as described herein, for example, backbone modifications having Formula I. In addition, the 2'-deoxy nucleotide shown 5' to the terminal modifications shown can be another modified or unmodified nucleotide or non-nucleotide described herein, for example modifications having any of Formulae I-VII or any combination thereof.

Figure 11 shows a non-limiting example of a strategy used to identify chemically modified siNA constructs of the invention that are nuclease resistance while preserving the ability to mediate RNAi activity. Chemical modifications are introduced into the siNA construct based on educated design parameters (e.g. introducing 2'-modifications, base modifications, backbone modifications, terminal cap modifications etc). The modified construct is tested in an appropriate system (e.g. human serum for nuclease resistance, shown, or an animal model for PK/delivery parameters). In parallel, the siNA construct is tested for RNAi activity, for example in a cell culture system such as a luciferase reporter assay). Lead siNA constructs are then identified which possess a particular characteristic while maintaining RNAi activity, and can be further modified and assayed once again. This same approach can be used to identify siNA-conjugate molecules with improved pharmacokinetic profiles, delivery, and RNAi activity.

Figure 12 shows non-limiting examples of phosphorylated siNA molecules of the invention, including linear and duplex constructs and asymmetric derivatives thereof.

Figure 13 shows non-limiting examples of chemically modified terminal phosphate groups of the invention.

Figure 14A shows a non-limiting example of methodology used to design self complementary DFO constructs utilizing palidrome and/or repeat nucleic acid sequences that are identified in a target nucleic acid sequence. (i) A palidrome or repeat sequence is identified in a nucleic acid target sequence. (ii) A sequence is designed that is complementary to the target nucleic acid sequence and the palidrome sequence. (iii) An inverse repeat sequence of the non-palidrome/repeat portion of the complementary sequence is appended to the 3'-end of the complementary sequence to generate a self complementary DFO molecule comprising sequence complementary to the nucleic acid target. (iv) The DFO molecule can self-assemble to form a double stranded oligonucleotide. **Figure 14B** shows a non-limiting representative example of a duplex forming oligonucleotide sequence. **Figure 14C** shows a non-limiting example of the self assembly schematic of a representative duplex forming oligonucleotide sequence. **Figure 14D** shows a non-limiting example of the self assembly schematic of a representative duplex forming oligonucleotide sequence followed by interaction with a target nucleic acid sequence resulting in modulation of gene expression.

Figure 15 shows a non-limiting example of the design of self complementary DFO constructs utilizing palindrome and/or repeat nucleic acid sequences that are incorporated into the DFO constructs that have sequence complementary to any target nucleic acid sequence of interest. Incorporation of these palindrome/repeat sequences allow the design of DFO constructs that form duplexes in which each strand is capable of mediating modulation of target gene expression, for example by RNAi. First, the target sequence is identified. A complementary sequence is then generated in which nucleotide or non-nucleotide modifications (shown as X or Y) are introduced into the complementary sequence that generate an artificial palindrome (shown as XXXYXXY in the Figure). An inverse repeat of the non-palindrome/repeat complementary sequence is appended to the 3'-end of the complementary sequence to generate a self complementary DFO comprising sequence complementary to the nucleic acid target. The DFO can self-assemble to form a double stranded oligonucleotide.

Figure 16 shows non-limiting examples of multifunctional siNA molecules of the invention comprising two separate polynucleotide sequences that are each capable of mediating RNAi directed cleavage of differing target nucleic acid sequences. **Figure 16A** shows a non-limiting example of a multifunctional siNA molecule having a first region that is complementary to a first target nucleic acid sequence (complementary region 1) and a second region that is complementary to a second target nucleic acid sequence (complementary region 2), wherein the first and second complementary regions are situated at the 3'-ends of each polynucleotide sequence in the multifunctional siNA. The dashed portions of each polynucleotide sequence of the multifunctional siNA construct have complementarity with regard to corresponding portions of the siNA duplex, but do not have complementarity to the target nucleic acid sequences. **Figure 16B** shows a non-limiting example of a multifunctional siNA molecule having a first region that is complementary to a first target nucleic acid sequence (complementary region 1) and a second region that is complementary to a second target nucleic acid sequence (complementary region 2), wherein the first and second complementary regions are situated at the 5'-ends of each polynucleotide sequence in the multifunctional siNA. The dashed portions of each polynucleotide sequence of the multifunctional siNA construct have complementarity with regard to corresponding portions of the siNA duplex, but do not have complementarity to the target nucleic acid sequences.

Figure 17 shows non-limiting examples of multifunctional siNA molecules of the invention comprising a single polynucleotide sequence comprising distinct regions that are each capable of mediating RNAi directed cleavage of differing target nucleic acid sequences. Figure 17A shows a non-limiting example of a multifunctional siNA molecule having a first region that is complementary to a first target nucleic acid sequence (complementary region 1) and a second region that is complementary to a second target nucleic acid sequence (complementary region 2), wherein the second complementary region is situated at the 3'-end of the polynucleotide sequence in the multifunctional siNA. The dashed portions of each polynucleotide sequence of the multifunctional siNA construct have complementarity with regard to corresponding portions of the siNA duplex, but do not have complementarity to the target nucleic acid sequences. Figure 17B shows a non-limiting example of a multifunctional siNA molecule having a first region that is complementary to a first target nucleic acid sequence (complementary region 1) and a second region that is complementary to a second target nucleic acid sequence (complementary region 2), wherein the first complementary region is situated at the 5'-end of the polynucleotide sequence in the multifunctional siNA. The dashed portions of each polynucleotide sequence of the multifunctional siNA construct have complementarity with regard to corresponding portions of the siNA duplex, but do not have complementarity to the target nucleic acid sequences. In one embodiment, these multifunctional siNA constructs are processed in vivo or in vitro to generate multifunctional siNA constructs as shown in Figure 16.

Figure 18 shows non-limiting examples of multifunctional siNA molecules of the invention comprising two separate polynucleotide sequences that are each capable of mediating RNAi directed cleavage of differing target nucleic acid sequences and wherein the multifunctional siNA construct further comprises a self complementary, palindrome, or repeat region, thus enabling shorter bifunctional siNA constructs that can mediate RNA interference against differing target nucleic acid sequences. Figure 18A shows a non-limiting example of a multifunctional siNA molecule having a first region that is complementary to a first target nucleic acid sequence (complementary region 1) and a second region that is complementary to a second target nucleic acid sequence (complementary region 2), wherein the first and second complementary regions are situated at the 3'-ends of each polynucleotide sequence in the multifunctional siNA, and

wherein the first and second complementary regions further comprise a self complementary, palindrome, or repeat region. The dashed portions of each polynucleotide sequence of the multifunctional siNA construct have complementarity with regard to corresponding portions of the siNA duplex, but do not have

5 complementarity to the target nucleic acid sequences. **Figure 18B** shows a non-limiting example of a multifunctional siNA molecule having a first region that is complementary to a first target nucleic acid sequence (complementary region 1) and a second region that is complementary to a second target nucleic acid sequence (complementary region 2), wherein the first and second complementary regions are situated at the 5'-ends of each

10 polynucleotide sequence in the multifunctional siNA, and wherein the first and second complementary regions further comprise a self complementary, palindrome, or repeat region. The dashed portions of each polynucleotide sequence of the multifunctional siNA construct have complementarity with regard to corresponding portions of the siNA duplex, but do not have complementarity to the target nucleic acid sequences.

15 **Figure 19** shows non-limiting examples of multifunctional siNA molecules of the invention comprising a single polynucleotide sequence comprising distinct regions that are each capable of mediating RNAi directed cleavage of differing target nucleic acid sequences and wherein the multifunctional siNA construct further comprises a self complementary, palindrome, or repeat region, thus enabling shorter bifunctional siNA

20 constructs that can mediate RNA interference against differing target nucleic acid sequences. **Figure 19A** shows a non-limiting example of a multifunctional siNA molecule having a first region that is complementary to a first target nucleic acid sequence (complementary region 1) and a second region that is complementary to a second target nucleic acid sequence (complementary region 2), wherein the second

25 complementary region is situated at the 3'-end of the polynucleotide sequence in the multifunctional siNA, and wherein the first and second complementary regions further comprise a self complementary, palindrome, or repeat region. The dashed portions of each polynucleotide sequence of the multifunctional siNA construct have complementarity with regard to corresponding portions of the siNA duplex, but do not

30 have complementarity to the target nucleic acid sequences. **Figure 19B** shows a non-limiting example of a multifunctional siNA molecule having a first region that is complementary to a first target nucleic acid sequence (complementary region 1) and a

second region that is complementary to a second target nucleic acid sequence (complementary region 2), wherein the first complementary region is situated at the 5'-end of the polynucleotide sequence in the multifunctional siNA, and wherein the first and second complementary regions further comprise a self complementary, palindrome, or repeat region. The dashed portions of each polynucleotide sequence of the multifunctional siNA construct have complementarity with regard to corresponding portions of the siNA duplex, but do not have complementarity to the target nucleic acid sequences. In one embodiment, these multifunctional siNA constructs are processed in vivo or in vitro to generate multifunctional siNA constructs as shown in Figure 18.

Figure 20 shows a non-limiting example of how multifunctional siNA molecules of the invention can target two separate target nucleic acid molecules, such as separate RNA molecules encoding differing proteins, for example, a cytokine and its corresponding receptor, differing viral strains, a virus and a cellular protein involved in viral infection or replication, or differing proteins involved in a common or divergent biologic pathway that is implicated in the maintenance of progression of disease. Each strand of the multifunctional siNA construct comprises a region having complementarity to separate target nucleic acid molecules. The multifunctional siNA molecule is designed such that each strand of the siNA can be utilized by the RISC complex to initiate RNA interference mediated cleavage of its corresponding target. These design parameters can include destabilization of each end of the siNA construct (see for example Schwarz *et al.*, 2003, *Cell*, 115, 199-208). Such destabilization can be accomplished for example by using guanosine-cytidine base pairs, alternate base pairs (e.g., wobbles), or destabilizing chemically modified nucleotides at terminal nucleotide positions as is known in the art.

Figure 21 shows a non-limiting example of how multifunctional siNA molecules of the invention can target two separate target nucleic acid sequences within the same target nucleic acid molecule, such as alternate coding regions of a RNA, coding and non-coding regions of a RNA, or alternate splice variant regions of a RNA. Each strand of the multifunctional siNA construct comprises a region having complementarity to the separate regions of the target nucleic acid molecule. The multifunctional siNA molecule is designed such that each strand of the siNA can be utilized by the RISC complex to

initiate RNA interference mediated cleavage of its corresponding target region. These design parameters can include destabilization of each end of the siNA construct (see for example Schwarz *et al.*, 2003, *Cell*, 115, 199-208). Such destabilization can be accomplished for example by using guanosine-cytidine base pairs, alternate base pairs (e.g., wobbles), or destabilizing chemically modified nucleotides at terminal nucleotide positions as is known in the art.

DETAILED DESCRIPTION OF THE INVENTION

Mechanism of Action of Nucleic Acid Molecules of the Invention

The discussion that follows discusses the proposed mechanism of RNA interference mediated by short interfering RNA as is presently known, and is not meant to be limiting and is not an admission of prior art. Applicant demonstrates herein that chemically-modified short interfering nucleic acids possess similar or improved capacity to mediate RNAi as do siRNA molecules and are expected to possess improved stability and activity *in vivo*; therefore, this discussion is not meant to be limiting only to siRNA and can be applied to siNA as a whole. By "improved capacity to mediate RNAi" or "improved RNAi activity" is meant to include RNAi activity measured *in vitro* and/or *in vivo* where the RNAi activity is a reflection of both the ability of the siNA to mediate RNAi and the stability of the siNAs of the invention. In this invention, the product of these activities can be increased *in vitro* and/or *in vivo* compared to an all RNA siRNA or a siNA containing a plurality of ribonucleotides. In some cases, the activity or stability of the siNA molecule can be decreased (i.e., less than ten-fold), but the overall activity of the siNA molecule is enhanced *in vitro* and/or *in vivo*.

RNA interference refers to the process of sequence specific post-transcriptional gene silencing in animals mediated by short interfering RNAs (siRNAs) (Fire *et al.*, 1998, *Nature*, 391, 806). The corresponding process in plants is commonly referred to as post-transcriptional gene silencing or RNA silencing and is also referred to as quelling in fungi. The process of post-transcriptional gene silencing is thought to be an evolutionarily-conserved cellular defense mechanism used to prevent the expression of foreign genes which is commonly shared by diverse flora and phyla (Fire *et al.*, 1999, *Trends Genet.*, 15, 358). Such protection from foreign gene expression may have

evolved in response to the production of double-stranded RNAs (dsRNAs) derived from viral infection or the random integration of transposon elements into a host genome via a cellular response that specifically destroys homologous single-stranded RNA or viral genomic RNA. The presence of dsRNA in cells triggers the RNAi response though a mechanism that has yet to be fully characterized. This mechanism appears to be different from the interferon response that results from dsRNA-mediated activation of protein kinase PKR and 2', 5'-oligoadenylate synthetase resulting in non-specific cleavage of mRNA by ribonuclease L.

The presence of long dsRNAs in cells stimulates the activity of a ribonuclease III enzyme referred to as Dicer. Dicer is involved in the processing of the dsRNA into short pieces of dsRNA known as short interfering RNAs (siRNAs) (Bernstein *et al.*, 2001, *Nature*, 409, 363). Short interfering RNAs derived from Dicer activity are typically about 21 to about 23 nucleotides in length and comprise about 19 base pair duplexes. Dicer has also been implicated in the excision of 21- and 22-nucleotide small temporal RNAs (stRNAs) from precursor RNA of conserved structure that are implicated in translational control (Hutvagner *et al.*, 2001, *Science*, 293, 834). The RNAi response also features an endonuclease complex containing a siRNA, commonly referred to as an RNA-induced silencing complex (RISC), which mediates cleavage of single-stranded RNA having sequence homologous to the siRNA. Cleavage of the target RNA takes place in the middle of the region complementary to the guide sequence of the siRNA duplex (Elbashir *et al.*, 2001, *Genes Dev.*, 15, 188). In addition, RNA interference can also involve small RNA (e.g., micro-RNA or miRNA) mediated gene silencing, presumably through cellular mechanisms that regulate chromatin structure and thereby prevent transcription of target gene sequences (see for example Allshire, 2002, *Science*, 297, 1818-1819; Volpe *et al.*, 2002, *Science*, 297, 1833-1837; Jenuwein, 2002, *Science*, 297, 2215-2218; and Hall *et al.*, 2002, *Science*, 297, 2232-2237). As such, siRNA molecules of the invention can be used to mediate gene silencing via interaction with RNA transcripts or alternately by interaction with particular gene sequences, wherein such interaction results in gene silencing either at the transcriptional level or post-transcriptional level.

RNAi has been studied in a variety of systems. Fire *et al.*, 1998, *Nature*, 391, 806, were the first to observe RNAi in *C. elegans*. Wianny and Goetz, 1999, *Nature Cell Biol.*, 2, 70, describe RNAi mediated by dsRNA in mouse embryos. Hammond *et al.*, 2000, *Nature*, 404, 293, describe RNAi in *Drosophila* cells transfected with dsRNA.

5 Elbashir *et al.*, 2001, *Nature*, 411, 494, describe RNAi induced by introduction of duplexes of synthetic 21-nucleotide RNAs in cultured mammalian cells including human embryonic kidney and HeLa cells. Recent work in *Drosophila* embryonic lysates has revealed certain requirements for siRNA length, structure, chemical composition, and sequence that are essential to mediate efficient RNAi activity. These studies have shown

10 that 21 nucleotide siRNA duplexes are most active when containing two 2-nucleotide 3'-terminal nucleotide overhangs. Furthermore, substitution of one or both siRNA strands with 2'-deoxy or 2'-O-methyl nucleotides abolishes RNAi activity, whereas substitution of 3'-terminal siRNA nucleotides with deoxy nucleotides was shown to be tolerated. Mismatch sequences in the center of the siRNA duplex were also shown to abolish RNAi

15 activity. In addition, these studies also indicate that the position of the cleavage site in the target RNA is defined by the 5'-end of the siRNA guide sequence rather than the 3'-end (Elbashir *et al.*, 2001, *EMBO J.*, 20, 6877). Other studies have indicated that a 5'-phosphate on the target-complementary strand of a siRNA duplex is required for siRNA activity and that ATP is utilized to maintain the 5'-phosphate moiety on the siRNA

20 (Nykanen *et al.*, 2001, *Cell*, 107, 309); however, siRNA molecules lacking a 5'-phosphate are active when introduced exogenously, suggesting that 5'-phosphorylation of siRNA constructs may occur *in vivo*.

Synthesis of Nucleic Acid Molecules

Synthesis of nucleic acids greater than 100 nucleotides in length is difficult using

25 automated methods, and the therapeutic cost of such molecules is prohibitive. In this invention, small nucleic acid motifs ("small" refers to nucleic acid motifs no more than 100 nucleotides in length, preferably no more than 80 nucleotides in length, and most preferably no more than 50 nucleotides in length; *e.g.*, individual siRNA oligonucleotide sequences or siRNA sequences synthesized in tandem) are preferably used for exogenous

30 delivery. The simple structure of these molecules increases the ability of the nucleic acid

to invade targeted regions of protein and/or RNA structure. Exemplary molecules of the instant invention are chemically synthesized, and others can similarly be synthesized.

Oligonucleotides (e.g., certain modified oligonucleotides or portions of oligonucleotides lacking ribonucleotides) are synthesized using protocols known in the art, for example as described in Caruthers *et al.*, 1992, *Methods in Enzymology* 211, 3-19, Thompson *et al.*, International PCT Publication No. WO 99/54459, Wincott *et al.*, 1995, *Nucleic Acids Res.* 23, 2677-2684, Wincott *et al.*, 1997, *Methods Mol. Bio.*, 74, 59, Brennan *et al.*, 1998, *Biotechnol Bioeng.*, 61, 33-45, and Brennan, U.S. Pat. No. 6,001,311. All of these references are incorporated herein by reference. The synthesis of oligonucleotides makes use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl at the 5'-end, and phosphoramidites at the 3'-end. In a non-limiting example, small scale syntheses are conducted on a 394 Applied Biosystems, Inc. synthesizer using a 0.2 μ mol scale protocol with a 2.5 min coupling step for 2'-O-methylated nucleotides and a 45 second coupling step for 2'-deoxy nucleotides or 2'-deoxy-2'-fluoro nucleotides. **Table III** outlines the amounts and the contact times of the reagents used in the synthesis cycle. Alternatively, syntheses at the 0.2 μ mol scale can be performed on a 96-well plate synthesizer, such as the instrument produced by Protogene (Palo Alto, CA) with minimal modification to the cycle. A 33-fold excess (60 μ L of 0.11 M = 6.6 μ mol) of 2'-O-methyl phosphoramidite and a 105-fold excess of S-ethyl tetrazole (60 μ L of 0.25 M = 15 μ mol) can be used in each coupling cycle of 2'-O-methyl residues relative to polymer-bound 5'-hydroxyl. A 22-fold excess (40 μ L of 0.11 M = 4.4 μ mol) of deoxy phosphoramidite and a 70-fold excess of S-ethyl tetrazole (40 μ L of 0.25 M = 10 μ mol) can be used in each coupling cycle of deoxy residues relative to polymer-bound 5'-hydroxyl. Average coupling yields on the 394 Applied Biosystems, Inc. synthesizer, determined by colorimetric quantitation of the trityl fractions, are typically 97.5-99%. Other oligonucleotide synthesis reagents for the 394 Applied Biosystems, Inc. synthesizer include the following: detritylation solution is 3% TCA in methylene chloride (ABI); capping is performed with 16% *N*-methyl imidazole in THF (ABI) and 10% acetic anhydride/10% 2,6-lutidine in THF (ABI); and oxidation solution is 16.9 mM I_2 , 49 mM pyridine, 9% water in THF (PerSeptive Biosystems, Inc.). Burdick & Jackson Synthesis Grade acetonitrile is used directly from the reagent bottle. S-Ethyltetrazole solution (0.25 M in acetonitrile) is made up from the solid obtained

from American International Chemical, Inc. Alternately, for the introduction of phosphorothioate linkages, Beaucage reagent (3H-1,2-Benzodithiol-3-one 1,1-dioxide, 0.05 M in acetonitrile) is used.

- Deprotection of the DNA-based oligonucleotides is performed as follows: the
- 5 polymer-bound trityl-on oligoribonucleotide is transferred to a 4 mL glass screw top vial and suspended in a solution of 40% aqueous methylamine (1 mL) at 65 °C for 10 minutes. After cooling to -20 °C, the supernatant is removed from the polymer support. The support is washed three times with 1.0 mL of EtOH:MeCN:H₂O/3:1:1, vortexed and the supernatant is then added to the first supernatant. The combined supernatants,
- 10 containing the oligoribonucleotide, are dried to a white powder.

- The method of synthesis used for RNA including certain siRNA molecules of the invention follows the procedure as described in Usman *et al.*, 1987, *J. Am. Chem. Soc.*, 109, 7845; Scaringe *et al.*, 1990, *Nucleic Acids Res.*, 18, 5433; and Wincott *et al.*, 1995, *Nucleic Acids Res.* 23, 2677-2684 Wincott *et al.*, 1997, *Methods Mol. Bio.*, 74, 59, and
- 15 makes use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl at the 5'-end, and phosphoramidites at the 3'-end. In a non-limiting example, small scale syntheses are conducted on a 394 Applied Biosystems, Inc. synthesizer using a 0.2 µmol scale protocol with a 7.5 min coupling step for alkylsilyl protected nucleotides and a 2.5 min coupling step for 2'-O-methylated nucleotides.
 - 20 **Table III** outlines the amounts and the contact times of the reagents used in the synthesis cycle. Alternatively, syntheses at the 0.2 µmol scale can be done on a 96-well plate synthesizer, such as the instrument produced by Protogene (Palo Alto, CA) with minimal modification to the cycle. A 33-fold excess (60 µL of 0.11 M = 6.6 µmol) of 2'-O-methyl phosphoramidite and a 75-fold excess of S-ethyl tetrazole (60 µL of 0.25 M = 15
 - 25 µmol) can be used in each coupling cycle of 2'-O-methyl residues relative to polymer-bound 5'-hydroxyl. A 66-fold excess (120 µL of 0.11 M = 13.2 µmol) of alkylsilyl (ribo) protected phosphoramidite and a 150-fold excess of S-ethyl tetrazole (120 µL of 0.25 M = 30 µmol) can be used in each coupling cycle of ribo residues relative to polymer-bound 5'-hydroxyl. Average coupling yields on the 394 Applied Biosystems, Inc.
 - 30 synthesizer, determined by colorimetric quantitation of the trityl fractions, are typically 97.5-99%. Other oligonucleotide synthesis reagents for the 394 Applied Biosystems,

- Inc. synthesizer include the following: detritylation solution is 3% TCA in methylene chloride (ABI); capping is performed with 16% *N*-methyl imidazole in THF (ABI) and 10% acetic anhydride/10% 2,6-lutidine in THF (ABI); oxidation solution is 16.9 mM I₂, 49 mM pyridine, 9% water in THF (PerSeptive Biosystems, Inc.).
- 5 Burdick & Jackson Synthesis Grade acetonitrile is used directly from the reagent bottle. S-Ethyltetrazole solution (0.25 M in acetonitrile) is made up from the solid obtained from American International Chemical, Inc. Alternately, for the introduction of phosphorothioate linkages, Beaucage reagent (3H-1,2-Benzodithiol-3-one 1,1-dioxide) 0.05 M in acetonitrile) is used.
- 10 Deprotection of the RNA is performed using either a two-pot or one-pot protocol. For the two-pot protocol, the polymer-bound trityl-on oligoribonucleotide is transferred to a 4 mL glass screw top vial and suspended in a solution of 40% aq. methylamine (1 mL) at 65 °C for 10 min. After cooling to -20 °C, the supernatant is removed from the polymer support. The support is washed three times with 1.0 mL of
- 15 EtOH:MeCN:H₂O/3:1:1, vortexed and the supernatant is then added to the first supernatant. The combined supernatants, containing the oligoribonucleotide, are dried to a white powder. The base deprotected oligoribonucleotide is resuspended in anhydrous TEA/HF/NMP solution (300 µL of a solution of 1.5 mL *N*-methylpyrrolidinone, 750 µL TEA and 1 mL TEA·3HF to provide a 1.4 M HF concentration) and heated to 65 °C.
- 20 After 1.5 h, the oligomer is quenched with 1.5 M NH₄HCO₃.

Alternatively, for the one-pot protocol, the polymer-bound trityl-on oligoribonucleotide is transferred to a 4 mL glass screw top vial and suspended in a solution of 33% ethanolic methylamine/DMSO: 1/1 (0.8 mL) at 65 °C for 15 minutes. The vial is brought to room temperature TEA·3HF (0.1 mL) is added and the vial is

25 heated at 65 °C for 15 minutes. The sample is cooled at -20 °C and then quenched with 1.5 M NH₄HCO₃.

For purification of the trityl-on oligomers, the quenched NH₄HCO₃ solution is loaded onto a C-18 containing cartridge that had been prewashed with acetonitrile followed by 50 mM TEAA. After washing the loaded cartridge with water, the RNA is

30 detritylated with 0.5% TFA for 13 minutes. The cartridge is then washed again with

water, salt exchanged with 1 M NaCl and washed with water again. The oligonucleotide is then eluted with 30% acetonitrile.

The average stepwise coupling yields are typically >98% (Wincott *et al.*, 1995 *Nucleic Acids Res.* 23, 2677-2684). Those of ordinary skill in the art will recognize that
 5 the scale of synthesis can be adapted to be larger or smaller than the example described above including but not limited to 96-well format.

Alternatively, the nucleic acid molecules of the present invention can be synthesized separately and joined together post-synthetically, for example, by ligation (Moore *et al.*, 1992, *Science* 256, 9923; Draper *et al.*, International PCT publication No.
 10 WO 93/23569; Shabarova *et al.*, 1991, *Nucleic Acids Research* 19, 4247; Bellon *et al.*, 1997, *Nucleosides & Nucleotides*, 16, 951; Bellon *et al.*, 1997, *Bioconjugate Chem.* 8, 204), or by hybridization following synthesis and/or deprotection.

The siNA molecules of the invention can also be synthesized via a tandem synthesis methodology as described in Example 1 herein, wherein both siNA strands are
 15 synthesized as a single contiguous oligonucleotide fragment or strand separated by a cleavable linker which is subsequently cleaved to provide separate siNA fragments or strands that hybridize and permit purification of the siNA duplex. The linker can be a polynucleotide linker or a non-nucleotide linker. The tandem synthesis of siNA as described herein can be readily adapted to both multiwell/multiplate synthesis platforms
 20 such as 96 well or similarly larger multi-well platforms. The tandem synthesis of siNA as described herein can also be readily adapted to large scale synthesis platforms employing batch reactors, synthesis columns and the like.

A siNA molecule can also be assembled from two distinct nucleic acid strands or fragments wherein one fragment includes the sense region and the second fragment
 25 includes the antisense region of the RNA molecule.

The nucleic acid molecules of the present invention can be modified extensively to enhance stability by modification with nuclease resistant groups, for example, 2'-amino, 2'-C-allyl, 2'-fluoro, 2'-O-methyl, 2'-H (for a review see Usman and Cedergren, 1992, *TIBS* 17, 34; Usman *et al.*, 1994, *Nucleic Acids Symp. Ser.* 31, 163). siNA constructs can
 30 be purified by gel electrophoresis using general methods or can be purified by high

pressure liquid chromatography (HPLC; see Wincott *et al.*, *supra*, the totality of which is hereby incorporated herein by reference) and re-suspended in water.

In another aspect of the invention, siNA molecules of the invention are expressed from transcription units inserted into DNA or RNA vectors. The recombinant vectors can be DNA plasmids or viral vectors. siNA expressing viral vectors can be constructed based on, but not limited to, adeno-associated virus, retrovirus, adenovirus, or alphavirus. The recombinant vectors capable of expressing the siNA molecules can be delivered as described herein, and persist in target cells. Alternatively, viral vectors can be used that provide for transient expression of siNA molecules.

10 Optimizing Activity of the nucleic acid molecule of the invention.

Chemically synthesizing nucleic acid molecules with modifications (base, sugar and/or phosphate) can prevent their degradation by serum ribonucleases, which can increase their potency (see *e.g.*, Eckstein *et al.*, International Publication No. WO 92/07065; Perrault *et al.*, 1990 *Nature* 344, 565; Pieken *et al.*, 1991, *Science* 253, 314; Usman and Cedergren, 1992, *Trends in Biochem. Sci.* 17, 334; Usman *et al.*, International Publication No. WO 93/15187; and Rossi *et al.*, International Publication No. WO 91/03162; Sproat, U.S. Pat. No. 5,334,711; Gold *et al.*, U.S. Pat. No. 6,300,074; and Burgin *et al.*, *supra*; all of which are incorporated by reference herein). All of the above references describe various chemical modifications that can be made to the base, phosphate and/or sugar moieties of the nucleic acid molecules described herein. Modifications that enhance their efficacy in cells, and removal of bases from nucleic acid molecules to shorten oligonucleotide synthesis times and reduce chemical requirements are desired.

There are several examples in the art describing sugar, base and phosphate modifications that can be introduced into nucleic acid molecules with significant enhancement in their nuclease stability and efficacy. For example, oligonucleotides are modified to enhance stability and/or enhance biological activity by modification with nuclease resistant groups, for example, 2'-amino, 2'-C-allyl, 2'-fluoro, 2'-O-methyl, 2'-O-allyl, 2'-H, nucleotide base modifications (for a review see Usman and Cedergren, 1992, *TIBS*. 17, 34; Usman *et al.*, 1994, *Nucleic Acids Symp. Ser.* 31, 163; Burgin *et al.*, 1996,

- Biochemistry*, 35, 14090). Sugar modification of nucleic acid molecules have been extensively described in the art (see Eckstein *et al.*, *International Publication* PCT No. WO 92/07065; Perrault *et al.* *Nature*, 1990, 344, 565-568; Pieken *et al.* *Science*, 1991, 253, 314-317; Usman and Cedergren, *Trends in Biochem. Sci.*, 1992, 17, 334-339;
- 5 Usman *et al.* *International Publication* PCT No. WO 93/15187; Sproat, *U.S. Pat. No.* 5,334,711 and Beigelman *et al.*, 1995, *J. Biol. Chem.*, 270, 25702; Beigelman *et al.*, *International PCT publication No.* WO 97/26270; Beigelman *et al.*, *U.S. Pat. No.* 5,716,824; Usman *et al.*, *U.S. Pat. No.* 5,627,053; Woolf *et al.*, *International PCT Publication No.* WO 98/13526; Thompson *et al.*, *USSN* 60/082,404 which was filed on
- 10 April 20, 1998; Karpeisky *et al.*, 1998, *Tetrahedron Lett.*, 39, 1131; Earnshaw and Gait, 1998, *Biopolymers (Nucleic Acid Sciences)*, 48, 39-55; Verma and Eckstein, 1998, *Annu. Rev. Biochem.*, 67, 99-134; and Burlina *et al.*, 1997, *Bioorg. Med. Chem.*, 5, 1999-2010; all of the references are hereby incorporated in their totality by reference herein). Such publications describe general methods and strategies to determine the location of
- 15 incorporation of sugar, base and/or phosphate modifications and the like into nucleic acid molecules without modulating catalysis, and are incorporated by reference herein. In view of such teachings, similar modifications can be used as described herein to modify the siNA nucleic acid molecules of the instant invention so long as the ability of siNA to promote RNAi in cells is not significantly inhibited.
- 20 While chemical modification of oligonucleotide internucleotide linkages with phosphorothioate, phosphorodithioate, and/or 5'-methylphosphonate linkages improves stability, excessive modifications can cause some toxicity or decreased activity. Therefore, when designing nucleic acid molecules, the amount of these internucleotide linkages should be minimized. The reduction in the concentration of these linkages
- 25 should lower toxicity, resulting in increased efficacy and higher specificity of these molecules.

Short interfering nucleic acid (siNA) molecules having chemical modifications that maintain or enhance activity are provided. Such a nucleic acid is also generally more resistant to nucleases than an unmodified nucleic acid. Accordingly, the *in vitro* and/or

30 *in vivo* activity should not be significantly lowered. In cases in which modulation is the goal, therapeutic nucleic acid molecules delivered exogenously should optimally be

- stable within cells until translation of the target RNA has been modulated long enough to reduce the levels of the undesirable protein. This period of time varies between hours to days depending upon the disease state. Improvements in the chemical synthesis of RNA and DNA (Wincott *et al.*, 1995, *Nucleic Acids Res.* 23, 2677; Caruthers *et al.*, 1992, *Methods in Enzymology* 211, 3-19 (incorporated by reference herein)) have expanded the ability to modify nucleic acid molecules by introducing nucleotide modifications to enhance their nuclease stability, as described above.

- In one embodiment, nucleic acid molecules of the invention include one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) G-clamp nucleotides. A G-clamp nucleotide is a modified cytosine analog wherein the modifications confer the ability to hydrogen bond both Watson-Crick and Hoogsteen faces of a complementary guanine within a duplex, see for example Lin and Matteucci, 1998, *J. Am. Chem. Soc.*, 120, 8531-8532. A single G-clamp analog substitution within an oligonucleotide can result in substantially enhanced helical thermal stability and mismatch discrimination when hybridized to complementary oligonucleotides. The inclusion of such nucleotides in nucleic acid molecules of the invention results in both enhanced affinity and specificity to nucleic acid targets, complementary sequences, or template strands. In another embodiment, nucleic acid molecules of the invention include one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) LNA "locked nucleic acid" nucleotides such as a 2', 4'-C methylene bicyclo nucleotide (see for example Wengel *et al.*, International PCT Publication No. WO 00/66604 and WO 99/14226).

- In another embodiment, the invention features conjugates and/or complexes of siRNA molecules of the invention. Such conjugates and/or complexes can be used to facilitate delivery of siRNA molecules into a biological system, such as a cell. The conjugates and complexes provided by the instant invention can impart therapeutic activity by transferring therapeutic compounds across cellular membranes, altering the pharmacokinetics, and/or modulating the localization of nucleic acid molecules of the invention. The present invention encompasses the design and synthesis of novel conjugates and complexes for the delivery of molecules, including, but not limited to, small molecules, lipids, cholesterol, phospholipids, nucleosides, nucleotides, nucleic acids, antibodies, toxins, negatively charged polymers and other polymers, for example

proteins, peptides, hormones, carbohydrates, polyethylene glycols, or polyamines, across cellular membranes. In general, the transporters described are designed to be used either individually or as part of a multi-component system, with or without degradable linkers. These compounds are expected to improve delivery and/or localization of nucleic acid molecules of the invention into a number of cell types originating from different tissues, in the presence or absence of serum (see Sullenger and Cech, U.S. Pat. No. 5,854,038). Conjugates of the molecules described herein can be attached to biologically active molecules via linkers that are biodegradable, such as biodegradable nucleic acid linker molecules.

10 The term "biodegradable linker" as used herein, refers to a nucleic acid or non-nucleic acid linker molecule that is designed as a biodegradable linker to connect one molecule to another molecule, for example, a biologically active molecule to a siNA molecule of the invention or the sense and antisense strands of a siNA molecule of the invention. The biodegradable linker is designed such that its stability can be modulated
15 for a particular purpose, such as delivery to a particular tissue or cell type. The stability of a nucleic acid-based biodegradable linker molecule can be modulated by using various chemistries, for example combinations of ribonucleotides, deoxyribonucleotides, and chemically-modified nucleotides, such as 2'-O-methyl, 2'-fluoro, 2'-amino, 2'-O-amino, 2'-C-allyl, 2'-O-allyl, and other 2'-modified or base modified nucleotides. The
20 biodegradable nucleic acid linker molecule can be a dimer, trimer, tetramer or longer nucleic acid molecule, for example, an oligonucleotide of about 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, or 20 nucleotides in length, or can comprise a single nucleotide with a phosphorus-based linkage, for example, a phosphoramidate or phosphodiester linkage. The biodegradable nucleic acid linker molecule can also
25 comprise nucleic acid backbone, nucleic acid sugar, or nucleic acid base modifications.

The term "biodegradable" as used herein, refers to degradation in a biological system, for example, enzymatic degradation or chemical degradation.

The term "biologically active molecule" as used herein refers to compounds or molecules that are capable of eliciting or modifying a biological response in a system.
30 Non-limiting examples of biologically active siNA molecules either alone or in combination with other molecules contemplated by the instant invention include

therapeutically active molecules such as antibodies, cholesterol, hormones, antivirals, peptides, proteins, chemotherapeutics, small molecules, vitamins, co-factors, nucleosides, nucleotides, oligonucleotides, enzymatic nucleic acids, antisense nucleic acids, triplex forming oligonucleotides, 2,5-A chimeras, siNA, dsRNA, allozymes, aptamers, decoys and analogs thereof. Biologically active molecules of the invention also include molecules capable of modulating the pharmacokinetics and/or pharmacodynamics of other biologically active molecules, for example, lipids and polymers such as polyamines, polyamides, polyethylene glycol and other polyethers.

The term "phospholipid" as used herein, refers to a hydrophobic molecule comprising at least one phosphorus group. For example, a phospholipid can comprise a phosphorus-containing group and saturated or unsaturated alkyl group, optionally substituted with OH, COOH, oxo, amine, or substituted or unsubstituted aryl groups.

Therapeutic nucleic acid molecules (e.g., siNA molecules) delivered exogenously optimally are stable within cells until reverse transcription of the RNA has been modulated long enough to reduce the levels of the RNA transcript. The nucleic acid molecules are resistant to nucleases in order to function as effective intracellular therapeutic agents. Improvements in the chemical synthesis of nucleic acid molecules described in the instant invention and in the art have expanded the ability to modify nucleic acid molecules by introducing nucleotide modifications to enhance their nuclease stability as described above.

In yet another embodiment, siNA molecules having chemical modifications that maintain or enhance enzymatic activity of proteins involved in RNAi are provided. Such nucleic acids are also generally more resistant to nucleases than unmodified nucleic acids. Thus, *in vitro* and/or *in vivo* the activity should not be significantly lowered.

Use of the nucleic acid-based molecules of the invention will lead to better treatments by affording the possibility of combination therapies (e.g., multiple siNA molecules targeted to different genes; nucleic acid molecules coupled with known small molecule modulators; or intermittent treatment with combinations of molecules, including different motifs and/or other chemical or biological molecules). The treatment of subjects with siNA molecules can also include combinations of different types of

nucleic acid molecules, such as enzymatic nucleic acid molecules (ribozymes),
allozymes, antisense, 2,5-A oligoadenylate, decoys, and aptamers.

In another aspect a siNA molecule of the invention comprises one or more 5'
and/or a 3'-cap structure, for example, on only the sense siNA strand, the antisense siNA
5 strand, or both siNA strands.

By "cap structure" is meant chemical modifications, which have been incorporated
at either terminus of the oligonucleotide (see, for example, Adamic *et al.*, U.S. Pat. No.
5,998,203, incorporated by reference herein). These terminal modifications protect the
nucleic acid molecule from exonuclease degradation, and may help in delivery and/or
10 localization within a cell. The cap may be present at the 5'-terminus (5'-cap) or at the 3'-
terminal (3'-cap) or may be present on both termini. In non-limiting examples, the 5'-cap
includes, but is not limited to, glyceryl, inverted deoxy abasic residue (moiety); 4',5'-
methylene nucleotide; 1-(beta-D-erythrofuranosyl) nucleotide, 4'-thio nucleotide;
carbocyclic nucleotide; 1,5-anhydrohexitol nucleotide; L-nucleotides; alpha-nucleotides;
15 modified base nucleotide; phosphorodithioate linkage; *threo*-pentofuranosyl nucleotide;
acyclic 3',4'-seco nucleotide; acyclic 3,4-dihydroxybutyl nucleotide; acyclic 3,5-
dihydroxypentyl nucleotide, 3'-3'-inverted nucleotide moiety; 3'-3'-inverted abasic
moiety; 3'-2'-inverted nucleotide moiety; 3'-2'-inverted abasic moiety; 1,4-butanediol
phosphate; 3'-phosphoramidate; hexylphosphate; aminohexyl phosphate; 3'-phosphate;
20 3'-phosphorothioate; phosphorodithioate; or bridging or non-bridging
methylphosphonate moiety. Non-limiting examples of cap moieties are shown in Figure
10.

Non-limiting examples of the 3'-cap include, but are not limited to, glyceryl,
inverted deoxy abasic residue (moiety), 4', 5'-methylene nucleotide; 1-(beta-D-
25 erythrofuranosyl) nucleotide; 4'-thio nucleotide, carbocyclic nucleotide; 5'-amino-alkyl
phosphate; 1,3-diamino-2-propyl phosphate; 3-aminopropyl phosphate; 6-aminohexyl
phosphate; 1,2-aminododecyl phosphate; hydroxypropyl phosphate; 1,5-anhydrohexitol
nucleotide; L-nucleotide; alpha-nucleotide; modified base nucleotide;
phosphorodithioate; *threo*-pentofuranosyl nucleotide; acyclic 3',4'-seco nucleotide; 3,4-
30 dihydroxybutyl nucleotide; 3,5-dihydroxypentyl nucleotide, 5'-5'-inverted nucleotide
moiety; 5'-5'-inverted abasic moiety; 5'-phosphoramidate; 5'-phosphorothioate; 1,4-

butanediol phosphate; 5'-amino; bridging and/or non-bridging 5'-phosphoramidate, phosphorothioate and/or phosphorodithioate, bridging or non bridging methylphosphonate and 5'-mercapto moieties (for more details see Beaucage and Iyer, 1993, *Tetrahedron* 49, 1925; incorporated by reference herein).

- 5 By the term "non-nucleotide" is meant any group or compound which can be incorporated into a nucleic acid chain in the place of one or more nucleotide units, including either sugar and/or phosphate substitutions, and allows the remaining bases to exhibit their enzymatic activity. The group or compound is abasic in that it does not contain a commonly recognized nucleotide base, such as adenosine, guanine, cytosine, uracil or thymine and therefore lacks a base at the 1'-position.

- 10 An "alkyl" group refers to a saturated aliphatic hydrocarbon, including straight-chain, branched-chain, and cyclic alkyl groups. Preferably, the alkyl group has 1 to 12 carbons. More preferably, it is a lower alkyl of from 1 to 7 carbons, more preferably 1 to 4 carbons. The alkyl group can be substituted or unsubstituted. When substituted
- 15 substituted group(s) is preferably, hydroxyl, cyano, alkoxy, =O, =S, NO₂ or N(CH₃)₂, amino, or SH. The term also includes alkenyl groups that are unsaturated hydrocarbon groups containing at least one carbon-carbon double bond, including straight-chain, branched-chain, and cyclic groups. Preferably, the alkenyl group has 1 to 12 carbons. More preferably, it is a lower alkenyl of from 1 to 7 carbons, more preferably 1 to 4
- 20 carbons. The alkenyl group may be substituted or unsubstituted. When substituted the substituted group(s) is preferably, hydroxyl, cyano, alkoxy, =O, =S, NO₂, halogen, N(CH₃)₂, amino, or SH. The term "alkyl" also includes alkynyl groups that have an unsaturated hydrocarbon group containing at least one carbon-carbon triple bond, including straight-chain, branched-chain, and cyclic groups. Preferably, the alkynyl
- 25 group has 1 to 12 carbons. More preferably, it is a lower alkynyl of from 1 to 7 carbons, more preferably 1 to 4 carbons. The alkynyl group may be substituted or unsubstituted. When substituted the substituted group(s) is preferably, hydroxyl, cyano, alkoxy, =O, =S, NO₂ or N(CH₃)₂, amino or SH.

- 30 Such alkyl groups can also include aryl, alkylaryl, carbocyclic aryl, heterocyclic aryl, amide and ester groups. An "aryl" group refers to an aromatic group that has at

least one ring having a conjugated pi electron system and includes carbocyclic aryl, heterocyclic aryl and biaryl groups, all of which may be optionally substituted. The preferred substituent(s) of aryl groups are halogen, trihalomethyl, hydroxyl, SH, OH, cyano, alkoxy, alkyl, alkenyl, alkynyl, and amino groups. An "alkylaryl" group refers to an alkyl group (as described above) covalently joined to an aryl group (as described above). Carbocyclic aryl groups are groups wherein the ring atoms on the aromatic ring are all carbon atoms. The carbon atoms are optionally substituted. Heterocyclic aryl groups are groups having from 1 to 3 heteroatoms as ring atoms in the aromatic ring and the remainder of the ring atoms are carbon atoms. Suitable heteroatoms include oxygen, sulfur, and nitrogen, and include furanyl, thienyl, pyridyl, pyrrolyl, N-lower alkyl pyrrolo, pyrimidyl, pyrazinyl, imidazolyl and the like, all optionally substituted. An "amide" refers to an -C(O)-NH-R, where R is either alkyl, aryl, alkylaryl or hydrogen. An "ester" refers to an -C(O)-OR', where R is either alkyl, aryl, alkylaryl or hydrogen.

By "nucleotide" as used herein is as recognized in the art to include natural bases (standard), and modified bases well known in the art. Such bases are generally located at the 1' position of a nucleotide sugar moiety. Nucleotides generally comprise a base, sugar and a phosphate group. The nucleotides can be unmodified or modified at the sugar, phosphate and/or base moiety, (also referred to interchangeably as nucleotide analogs, modified nucleotides, non-natural nucleotides, non-standard nucleotides and other; see, for example, Usman and McSwiggen, *supra*; Eckstein *et al.*, International PCT Publication No. WO 92/07065; Usman *et al.*, International PCT Publication No. WO 93/15187; Uhlman & Peyman, *supra*, all are hereby incorporated by reference herein). There are several examples of modified nucleic acid bases known in the art as summarized by Limbach *et al.*, 1994, *Nucleic Acids Res.* 22, 2183. Some of the non-limiting examples of base modifications that can be introduced into nucleic acid molecules include, inosine, purine, pyridin-4-one, pyridin-2-one, phenyl, pseudouracil, 2, 4, 6-trimethoxy benzene, 3-methyl uracil, dihydrouridine, naphthyl, aminophenyl, 5-alkylcytidines (e.g., 5-methylcytidine), 5-alkyluridines (e.g., ribothymidine), 5-halouridine (e.g., 5-bromouridine) or 6-azapyrimidines or 6-alkylpyrimidines (e.g. 6-methyluridine), propyne, and others (Burgin *et al.*, 1996, *Biochemistry*, 35, 14090; Uhlman & Peyman, *supra*). By "modified bases" in this aspect is meant nucleotide bases other than adenine, guanine, cytosine and uracil at 1' position or their equivalents.

In one embodiment, the invention features modified siNA molecules, with phosphate backbone modifications comprising one or more phosphorothioate, phosphorodithioate, methylphosphonate, phosphotriester, morpholino, amide carbamate, carboxymethyl, acetamidate, polyamide, sulfonate, sulfonamide, sulfamate, formacetal, thioformacetal, and/or alkylsilyl, substitutions. For a review of oligonucleotide backbone modifications, see Hunziker and Leumann, 1995, *Nucleic Acid Analogues: Synthesis and Properties*, in *Modern Synthetic Methods*, VCH, 331-417, and Mesmaeker *et al.*, 1994, *Novel Backbone Replacements for Oligonucleotides*, in *Carbohydrate Modifications in Antisense Research*, ACS, 24-39.

By "abasic" is meant sugar moieties lacking a base or having other chemical groups in place of a base at the 1' position, see for example Adamic *et al.*, U.S. Pat. No. 5,998,203.

By "unmodified nucleoside" is meant one of the bases adenine, cytosine, guanine, thymine, or uracil joined to the 1' carbon of β -D-ribo-furanose.

By "modified nucleoside" is meant any nucleotide base which contains a modification in the chemical structure of an unmodified nucleotide base, sugar and/or phosphate. Non-limiting examples of modified nucleotides are shown by Formulae I-VII and/or other modifications described herein.

In connection with 2'-modified nucleotides as described for the present invention, by "amino" is meant 2'-NH₂ or 2'-O- NH₂, which can be modified or unmodified. Such modified groups are described, for example, in Eckstein *et al.*, U.S. Pat. No. 5,672,695 and Matulic-Adamic *et al.*, U.S. Pat. No. 6,248,878, which are both incorporated by reference in their entireties.

Various modifications to nucleic acid siNA structure can be made to enhance the utility of these molecules. Such modifications will enhance shelf-life, half-life *in vitro*, stability, and ease of introduction of such oligonucleotides to the target site, *e.g.*, to enhance penetration of cellular membranes, and confer the ability to recognize and bind to targeted cells.

Administration of Nucleic Acid Molecules

A siRNA molecule of the invention can be adapted for use to prevent or treat diseases, traits, disorders, and/or conditions described herein or otherwise known in the art to be related to gene expression, and/or any other trait, disease, disorder or condition that is related to or will respond to the levels of a target polynucleotide in a cell or tissue,

5 alone or in combination with other therapies. For example, a siRNA molecule can comprise a delivery vehicle, including liposomes, for administration to a subject, carriers and diluents and their salts, and/or can be present in pharmaceutically acceptable formulations. Methods for the delivery of nucleic acid molecules are described in Akhtar *et al.*, 1992, *Trends Cell Bio.*, 2, 139; *Delivery Strategies for Antisense*

10 *Oligonucleotide Therapeutics*, ed. Akhtar, 1995, Maurer *et al.*, 1999, *Mol. Membr. Biol.*, 16, 129-140; Hofland and Huang, 1999, *Handb. Exp. Pharmacol.*, 137, 165-192; and Lee *et al.*, 2000, *ACS Symp. Ser.*, 752, 184-192, all of which are incorporated herein by reference. Beigelman *et al.*, U.S. Pat. No. 6,395,713 and Sullivan *et al.*, PCT WO 94/02595 further describe the general methods for delivery of nucleic acid molecules.

15 These protocols can be utilized for the delivery of virtually any nucleic acid molecule. Nucleic acid molecules can be administered to cells by a variety of methods known to those of skill in the art, including, but not restricted to, encapsulation in liposomes, by iontophoresis, or by incorporation into other vehicles, such as biodegradable polymers, hydrogels, cyclodextrins (see for example Gonzalez *et al.*, 1999, *Bioconjugate Chem.*,

20 10, 1068-1074; Wang *et al.*, International PCT publication Nos. WO 03/47518 and WO 03/46185), poly(lactic-co-glycolic)acid (PLGA) and PLGA microspheres (see for example US Patent 6,447,796 and US Patent Application Publication No. US 2002130430), biodegradable nanocapsules, and bioadhesive microspheres, or by proteinaceous vectors (O'Hare and Normand, International PCT Publication No. WO

25 00/53722). Alternatively, the nucleic acid/vehicle combination is locally delivered by direct injection or by use of an infusion pump. Direct injection of the nucleic acid molecules of the invention, whether subcutaneous, intramuscular, or intradermal, can take place using standard needle and syringe methodologies, or by needle-free technologies such as those described in Conry *et al.*, 1999, *Clin. Cancer Res.*, 5, 2330-

30 2337 and Barry *et al.*, International PCT Publication No. WO 99/31262. The molecules of the instant invention can be used as pharmaceutical agents. Pharmaceutical agents

prevent, modulate the occurrence, or treat (alleviate a symptom to some extent, preferably all of the symptoms) of a disease state in a subject.

In another embodiment, the nucleic acid molecules of the invention can also be formulated or complexed with polyethyleneimine and derivatives thereof, such as polyethyleneimine-polyethyleneglycol-N-acetylgalactosamine (PEI-PEG-GAL) or polyethyleneimine-polyethyleneglycol-tri-N-acetylgalactosamine (PEI-PEG-triGAL) derivatives. In one embodiment, the nucleic acid molecules of the invention are formulated as described in United States Patent Application Publication No. 20030077829, incorporated by reference herein in its entirety.

In one embodiment, a siNA molecule of the invention is complexed with membrane disruptive agents such as those described in U.S. Patent Application Publication No. 20010007666, incorporated by reference herein in its entirety including the drawings. In another embodiment, the membrane disruptive agent or agents and the siNA molecule are also complexed with a cationic lipid or helper lipid molecule, such as those lipids described in U.S. Patent No. 6,235,310, incorporated by reference herein in its entirety including the drawings.

In one embodiment, a siNA molecule of the invention is complexed with delivery systems as described in U.S. Patent Application Publication No. 2003077829 and International PCT Publication Nos. WO 00/03683 and WO 02/087541, all incorporated by reference herein in their entirety including the drawings.

In one embodiment, the nucleic acid molecules of the invention are administered via pulmonary delivery, such as by inhalation of an aerosol or spray dried formulation administered by an inhalation device or nebulizer, providing rapid local uptake of the nucleic acid molecules into relevant pulmonary tissues. Solid particulate compositions containing respirable dry particles of micronized nucleic acid compositions can be prepared by grinding dried or lyophilized nucleic acid compositions, and then passing the micronized composition through, for example, a 400 mesh screen to break up or separate out large agglomerates. A solid particulate composition comprising the nucleic acid compositions of the invention can optionally contain a dispersant which serves to facilitate the formation of an aerosol as well as other therapeutic compounds. A suitable

dispersant is lactose, which can be blended with the nucleic acid compound in any suitable ratio, such as a 1 to 1 ratio by weight.

Aerosols of liquid particles comprising a nucleic acid composition of the invention can be produced by any suitable means, such as with a nebulizer (see for example US 4,501,729). Nebulizers are commercially available devices which transform solutions or suspensions of an active ingredient into a therapeutic aerosol mist either by means of acceleration of a compressed gas, typically air or oxygen, through a narrow venturi orifice or by means of ultrasonic agitation. Suitable formulations for use in nebulizers comprise the active ingredient in a liquid carrier in an amount of up to 40% w/w preferably less than 20% w/w of the formulation. The carrier is typically water or a dilute aqueous alcoholic solution, preferably made isotonic with body fluids by the addition of, for example, sodium chloride or other suitable salts. Optional additives include preservatives if the formulation is not prepared sterile, for example, methyl hydroxybenzoate, anti-oxidants, flavorings, volatile oils, buffering agents and emulsifiers and other formulation surfactants. The aerosols of solid particles comprising the active composition and surfactant can likewise be produced with any solid particulate aerosol generator. Aerosol generators for administering solid particulate therapeutics to a subject produce particles which are respirable, as explained above, and generate a volume of aerosol containing a predetermined metered dose of a therapeutic composition at a rate suitable for human administration. One illustrative type of solid particulate aerosol generator is an insufflator. Suitable formulations for administration by insufflation include finely comminuted powders which can be delivered by means of an insufflator. In the insufflator, the powder, e.g., a metered dose thereof effective to carry out the treatments described herein, is contained in capsules or cartridges, typically made of gelatin or plastic, which are either pierced or opened in situ and the powder delivered by air drawn through the device upon inhalation or by means of a manually-operated pump. The powder employed in the insufflator consists either solely of the active ingredient or of a powder blend comprising the active ingredient, a suitable powder diluent, such as lactose, and an optional surfactant. The active ingredient typically comprises from 0.1 to 100 w/w of the formulation. A second type of illustrative aerosol generator comprises a metered dose inhaler. Metered dose inhalers are pressurized aerosol dispensers, typically containing a suspension or solution formulation of the active ingredient in a liquefied

propellant. During use these devices discharge the formulation through a valve adapted to deliver a metered volume to produce a fine particle spray containing the active ingredient. Suitable propellants include certain chlorofluorocarbon compounds, for example, dichlorodifluoromethane, trichlorofluoromethane, dichlorotetrafluoroethane and mixtures thereof. The formulation can additionally contain one or more co-solvents, for example, ethanol, emulsifiers and other formulation surfactants, such as oleic acid or sorbitan trioleate, anti-oxidants and suitable flavoring agents. Other methods for pulmonary delivery are described in, for example US Patent Application No. 20040037780, and US Patent Nos. 6,592,904; 6,582,728; 6,565,885.

10 In one embodiment, the invention features the use of methods to deliver the nucleic acid molecules of the instant invention to the central nervous system and/or peripheral nervous system. Experiments have demonstrated the efficient *in vivo* uptake of nucleic acids by neurons. As an example of local administration of nucleic acids to nerve cells, Sommer *et al.*, 1998, *Antisense Nuc. Acid Drug Dev.*, 8, 75, describe a study in which a
15 15mer phosphorothioate antisense nucleic acid molecule to c-fos is administered to rats via microinjection into the brain. Antisense molecules labeled with tetramethylrhodamine-isothiocyanate (TRITC) or fluorescein isothiocyanate (FITC) were taken up by exclusively by neurons thirty minutes post-injection. A diffuse cytoplasmic staining and nuclear staining was observed in these cells. As an example of systemic
20 administration of nucleic acid to nerve cells, Epa *et al.*, 2000, *Antisense Nuc. Acid Drug Dev.*, 10, 469, describe an *in vivo* mouse study in which beta-cyclodextrin-adamantane-oligonucleotide conjugates were used to target the p75 neurotrophin receptor in neuronally differentiated PC12 cells. Following a two week course of IP administration, pronounced uptake of p75 neurotrophin receptor antisense was observed in dorsal root
25 ganglion (DRG) cells. In addition, a marked and consistent down-regulation of p75 was observed in DRG neurons. Additional approaches to the targeting of nucleic acid to neurons are described in Broaddus *et al.*, 1998, *J. Neurosurg.*, 88(4), 734; Karle *et al.*, 1997, *Eur. J. Pharmacol.*, 340(2/3), 153; Bannai *et al.*, 1998, *Brain Research*, 784(1,2), 304; Rajakumar *et al.*, 1997, *Synapse*, 26(3), 199; Wu-pong *et al.*, 1999, *BioPharm*,
30 12(1), 32; Bannai *et al.*, 1998, *Brain Res. Protoc.*, 3(1), 83; Simantov *et al.*, 1996, *Neuroscience*, 74(1), 39. Nucleic acid molecules of the invention are therefore amenable to delivery to and uptake by cells that express repeat expansion allelic variants for

modulation of RE gene expression. The delivery of nucleic acid molecules of the invention, targeting RE is provided by a variety of different strategies. Traditional approaches to CNS delivery that can be used include, but are not limited to, intrathecal and intracerebroventricular administration, implantation of catheters and pumps, direct
5 injection or perfusion at the site of injury or lesion, injection into the brain arterial system, or by chemical or osmotic opening of the blood-brain barrier. Other approaches can include the use of various transport and carrier systems, for example though the use of conjugates and biodegradable polymers. Furthermore, gene therapy approaches, for example as described in Kaplitt *et al.*, US 6,180,613 and Davidson, WO 04/013280, can
10 be used to express nucleic acid molecules in the CNS.

In one embodiment, nucleic acid molecules of the invention are administered to the central nervous system (CNS) or peripheral nervous system (PNS). Experiments have demonstrated the efficient *in vivo* uptake of nucleic acids by neurons. As an example of local administration of nucleic acids to nerve cells, Sommer *et al.*, 1998, *Antisense Nuc.*
15 *Acid Drug Dev.*, 8, 75, describe a study in which a 15mer phosphorothioate antisense nucleic acid molecule to c-fos is administered to rats via microinjection into the brain. Antisense molecules labeled with tetramethylrhodamine-isothiocyanate (TRITC) or fluorescein isothiocyanate (FITC) were taken up by exclusively by neurons thirty minutes post-injection. A diffuse cytoplasmic staining and nuclear staining was
20 observed in these cells. As an example of systemic administration of nucleic acid to nerve cells, Epa *et al.*, 2000, *Antisense Nuc. Acid Drug Dev.*, 10, 469, describe an *in vivo* mouse study in which beta-cyclodextrin-adamantane-oligonucleotide conjugates were used to target the p75 neurotrophin receptor in neuronally differentiated PC12 cells. Following a two week course of IP administration, pronounced uptake of p75
25 neurotrophin receptor antisense was observed in dorsal root ganglion (DRG) cells. In addition, a marked and consistent down-regulation of p75 was observed in DRG neurons. Additional approaches to the targeting of nucleic acid to neurons are described in Broaddus *et al.*, 1998, *J. Neurosurg.*, 88(4), 734; Karle *et al.*, 1997, *Eur. J. Pharmacol.*, 340(2/3), 153; Bannai *et al.*, 1998, *Brain Research*, 784(1,2), 304;
30 Rajakumar *et al.*, 1997, *Synapse*, 26(3), 199; Wu-pong *et al.*, 1999, *BioPharm*, 12(1), 32; Bannai *et al.*, 1998, *Brain Res. Protoc.*, 3(1), 83; Simantov *et al.*, 1996, *Neuroscience*,

74(1), 39. Nucleic acid molecules of the invention are therefore amenable to delivery to and uptake by cells in the CNS and/or PNS.

The delivery of nucleic acid molecules of the invention to the CNS is provided by a variety of different strategies. Traditional approaches to CNS delivery that can be used include, but are not limited to, intrathecal and intracerebroventricular administration, implantation of catheters and pumps, direct injection or perfusion at the site of injury or lesion, injection into the brain arterial system, or by chemical or osmotic opening of the blood-brain barrier. Other approaches can include the use of various transport and carrier systems, for example though the use of conjugates and biodegradable polymers. Furthermore, gene therapy approaches, for example as described in Kaplitt *et al.*, US 6,180,613 and Davidson, WO 04/013280, can be used to express nucleic acid molecules in the CNS.

In one embodiment, delivery systems of the invention include, for example, aqueous and nonaqueous gels, creams, multiple emulsions, microemulsions, liposomes, ointments, aqueous and nonaqueous solutions, lotions, aerosols, hydrocarbon bases and powders, and can contain excipients such as solubilizers, permeation enhancers (e.g., fatty acids, fatty acid esters, fatty alcohols and amino acids), and hydrophilic polymers (e.g., polycarbophil and polyvinylpyrrolidone). In one embodiment, the pharmaceutically acceptable carrier is a liposome or a transdermal enhancer. Examples of liposomes which can be used in this invention include the following: (1) Cellfectin, 1:1.5 (M/M) liposome formulation of the cationic lipid N,N,N,N-tetramethyl-N,N,N,N-tetrapalmityl-spermine and dioleoyl phosphatidylethanolamine (DOPE) (GIBCO BRL); (2) Cytofectin GSV, 2:1 (M/M) liposome formulation of a cationic lipid and DOPE (Glen Research); (3) DOTAP (N-[1-(2,3-dioleoyloxy)-N,N,N-tri-methyl-ammoniummethylsulfate) (Boehringer Mannheim); and (4) Lipofectamine, 3:1 (M/M) liposome formulation of the polycationic lipid DOSPA and the neutral lipid DOPE (GIBCO BRL).

In one embodiment, delivery systems of the invention include patches, tablets, suppositories, pessaries, gels and creams, and can contain excipients such as solubilizers and enhancers (e.g., propylene glycol, bile salts and amino acids), and other vehicles (e.g., polyethylene glycol, fatty acid esters and derivatives, and hydrophilic polymers such as hydroxypropylmethylcellulose and hyaluronic acid).

In one embodiment, siNA molecules of the invention are formulated or complexed with polyethylenimine (e.g., linear or branched PEI) and/or polyethylenimine derivatives, including for example grafted PEIs such as galactose PEI, cholesterol PEI, antibody derivatized PEI, and polyethylene glycol PEI (PEG-PEI) derivatives thereof

5 (see for example Ogris *et al.*, 2001, *AAPA PharmSci*, 3, 1-11; Furgeson *et al.*, 2003, Bioconjugate Chem., 14, 840-847; Kunath *et al.*, 2002, Pharmaceutical Research, 19, 810-817; Choi *et al.*, 2001, Bull. Korean Chem. Soc., 22, 46-52; Bettinger *et al.*, 1999, Bioconjugate Chem., 10, 558-561; Peterson *et al.*, 2002, Bioconjugate Chem., 13, 845-854; Erbacher *et al.*, 1999, Journal of Gene Medicine Preprint, 1, 1-18; Godbey *et al.*,

10 1999., PNAS USA, 96, 5177-5181; Godbey *et al.*, 1999, Journal of Controlled Release, 60, 149-160; Diebold *et al.*, 1999, Journal of Biological Chemistry, 274, 19087-19094; Thomas and Klibanov, 2002, PNAS USA, 99, 14640-14645; and Sagara, US 6,586,524, incorporated by reference herein.

In one embodiment, a siNA molecule of the invention comprises a bioconjugate,

15 for example a nucleic acid conjugate as described in Vargeese *et al.*, USSN 10/427,160, filed April 30, 2003; US 6,528,631; US 6,335,434; US 6, 235,886; US 6,153,737; US 5,214,136; US 5,138,045, all incorporated by reference herein.

Thus, the invention features a pharmaceutical composition comprising one or more nucleic acid(s) of the invention in an acceptable carrier, such as a stabilizer, buffer, and the like. The polynucleotides of the invention can be administered (e.g., RNA, DNA or

20 protein) and introduced to a subject by any standard means, with or without stabilizers, buffers, and the like, to form a pharmaceutical composition. When it is desired to use a liposome delivery mechanism, standard protocols for formation of liposomes can be followed. The compositions of the present invention can also be formulated and used as

25 creams, gels, sprays, oils and other suitable compositions for topical, dermal, or transdermal administration as is known in the art.

The present invention also includes pharmaceutically acceptable formulations of the compounds described. These formulations include salts of the above compounds, e.g., acid addition salts, for example, salts of hydrochloric, hydrobromic, acetic acid, and

30 benzene sulfonic acid.

A pharmacological composition or formulation refers to a composition or formulation in a form suitable for administration, *e.g.*, systemic or local administration, into a cell or subject, including for example a human. Suitable forms, in part, depend upon the use or the route of entry, for example oral, transdermal, or by injection. Such forms should not prevent the composition or formulation from reaching a target cell (*i.e.*, a cell to which the negatively charged nucleic acid is desirable for delivery). For example, pharmacological compositions injected into the blood stream should be soluble. Other factors are known in the art, and include considerations such as toxicity and forms that prevent the composition or formulation from exerting its effect.

In one embodiment, siNA molecules of the invention are administered to a subject by systemic administration in a pharmaceutically acceptable composition or formulation. By "systemic administration" is meant *in vivo* systemic absorption or accumulation of drugs in the blood stream followed by distribution throughout the entire body. Administration routes that lead to systemic absorption include, without limitation: intravenous, subcutaneous, intraperitoneal, inhalation, oral, intrapulmonary and intramuscular. Each of these administration routes exposes the siNA molecules of the invention to an accessible diseased tissue. The rate of entry of a drug into the circulation has been shown to be a function of molecular weight or size. The use of a liposome or other drug carrier comprising the compounds of the instant invention can potentially localize the drug, for example, in certain tissue types, such as the tissues of the reticular endothelial system (RES). A liposome formulation that can facilitate the association of drug with the surface of cells, such as, lymphocytes and macrophages is also useful. This approach can provide enhanced delivery of the drug to target cells by taking advantage of the specificity of macrophage and lymphocyte immune recognition of abnormal cells, such as cancer cells.

By "pharmaceutically acceptable formulation" or "pharmaceutically acceptable composition" is meant, a composition or formulation that allows for the effective distribution of the nucleic acid molecules of the instant invention in the physical location most suitable for their desired activity. Non-limiting examples of agents suitable for formulation with the nucleic acid molecules of the instant invention include: P-glycoprotein inhibitors (such as Pluronic P85);, biodegradable polymers, such as poly

(DL-lactide-coglycolide) microspheres for sustained release delivery (Emerich, DF *et al.*, 1999, *Cell Transplant*, 8, 47-58); and loaded nanoparticles, such as those made of polybutylcyanoacrylate. Other non-limiting examples of delivery strategies for the nucleic acid molecules of the instant invention include material described in Boado *et al.*, 1998, *J. Pharm. Sci.*, 87, 1308-1315; Tyler *et al.*, 1999, *FEBS Lett.*, 421, 280-284; Pardridge *et al.*, 1995, *PNAS USA*, 92, 5592-5596; Boado, 1995, *Adv. Drug Delivery Rev.*, 15, 73-107; Aldrian-Herrada *et al.*, 1998, *Nucleic Acids Res.*, 26, 4910-4916; and Tyler *et al.*, 1999, *PNAS USA*, 96, 7053-7058.

The invention also features the use of the composition comprising surface-modified liposomes containing poly (ethylene glycol) lipids (PEG-modified, or long-circulating liposomes or stealth liposomes). These formulations offer a method for increasing the accumulation of drugs in target tissues. This class of drug carriers resists opsonization and elimination by the mononuclear phagocytic system (MPS or RES), thereby enabling longer blood circulation times and enhanced tissue exposure for the encapsulated drug (Lasic *et al. Chem. Rev.* 1995, 95, 2601-2627; Ishiwata *et al., Chem. Pharm. Bull.* 1995, 43, 1005-1011). Such liposomes have been shown to accumulate selectively in tumors, presumably by extravasation and capture in the neovascularized target tissues (Lasic *et al., Science* 1995, 267, 1275-1276; Oku *et al.*, 1995, *Biochim. Biophys. Acta*, 1238, 86-90). The long-circulating liposomes enhance the pharmacokinetics and pharmacodynamics of DNA and RNA, particularly compared to conventional cationic liposomes which are known to accumulate in tissues of the MPS (Liu *et al., J. Biol. Chem.* 1995, 42, 24864-24870; Choi *et al.*, International PCT Publication No. WO 96/10391; Ansell *et al.*, International PCT Publication No. WO 96/10390; Holland *et al.*, International PCT Publication No. WO 96/10392). Long-circulating liposomes are also likely to protect drugs from nuclease degradation to a greater extent compared to cationic liposomes, based on their ability to avoid accumulation in metabolically aggressive MPS tissues such as the liver and spleen.

The present invention also includes compositions prepared for storage or administration that include a pharmaceutically effective amount of the desired compounds in a pharmaceutically acceptable carrier or diluent. Acceptable carriers or diluents for therapeutic use are well known in the pharmaceutical art, and are described,

for example, in *Remington's Pharmaceutical Sciences*, Mack Publishing Co. (A.R. Gennaro edit. 1985), hereby incorporated by reference herein. For example, preservatives, stabilizers, dyes and flavoring agents can be provided. These include sodium benzoate, sorbic acid and esters of *p*-hydroxybenzoic acid. In addition, 5 antioxidants and suspending agents can be used.

A pharmaceutically effective dose is that dose required to prevent, inhibit the occurrence, or treat (alleviate a symptom to some extent, preferably all of the symptoms) of a disease state. The pharmaceutically effective dose depends on the type of disease, the composition used, the route of administration, the type of mammal being treated, the 10 physical characteristics of the specific mammal under consideration, concurrent medication, and other factors that those skilled in the medical arts will recognize. Generally, an amount between 0.1 mg/kg and 100 mg/kg body weight/day of active ingredients is administered dependent upon potency of the negatively charged polymer.

The nucleic acid molecules of the invention and formulations thereof can be 15 administered orally, topically, parenterally, by inhalation or spray, or rectally in dosage unit formulations containing conventional non-toxic pharmaceutically acceptable carriers, adjuvants and/or vehicles. The term parenteral as used herein includes percutaneous, subcutaneous, intravascular (*e.g.*, intravenous), intramuscular, or intrathecal injection or infusion techniques and the like. In addition, there is provided a 20 pharmaceutical formulation comprising a nucleic acid molecule of the invention and a pharmaceutically acceptable carrier. One or more nucleic acid molecules of the invention can be present in association with one or more non-toxic pharmaceutically acceptable carriers and/or diluents and/or adjuvants, and if desired other active ingredients. The pharmaceutical compositions containing nucleic acid molecules of the 25 invention can be in a form suitable for oral use, for example, as tablets, troches, lozenges, aqueous or oily suspensions, dispersible powders or granules, emulsion, hard or soft capsules, or syrups or elixirs.

Compositions intended for oral use can be prepared according to any method known to the art for the manufacture of pharmaceutical compositions and such 30 compositions can contain one or more such sweetening agents, flavoring agents, coloring agents or preservative agents in order to provide pharmaceutically elegant and palatable

preparations. Tablets contain the active ingredient in admixture with non-toxic pharmaceutically acceptable excipients that are suitable for the manufacture of tablets. These excipients can be, for example, inert diluents; such as calcium carbonate, sodium carbonate, lactose, calcium phosphate or sodium phosphate; granulating and
5 disintegrating agents, for example, corn starch, or alginic acid; binding agents, for example starch, gelatin or acacia; and lubricating agents, for example magnesium stearate, stearic acid or talc. The tablets can be uncoated or they can be coated by known techniques. In some cases such coatings can be prepared by known techniques to delay disintegration and absorption in the gastrointestinal tract and thereby provide a sustained
10 action over a longer period. For example, a time delay material such as glyceryl monostearate or glyceryl distearate can be employed.

Formulations for oral use can also be presented as hard gelatin capsules wherein the active ingredient is mixed with an inert solid diluent, for example, calcium carbonate, calcium phosphate or kaolin, or as soft gelatin capsules wherein the active ingredient is
15 mixed with water or an oil medium, for example peanut oil, liquid paraffin or olive oil.

Aqueous suspensions contain the active materials in a mixture with excipients suitable for the manufacture of aqueous suspensions. Such excipients are suspending agents, for example sodium carboxymethylcellulose, methylcellulose, hydropropyl-methylcellulose, sodium alginate, polyvinylpyrrolidone, gum tragacanth and gum acacia;
20 dispersing or wetting agents can be a naturally-occurring phosphatide, for example, lecithin, or condensation products of an alkylene oxide with fatty acids, for example polyoxyethylene stearate, or condensation products of ethylene oxide with long chain aliphatic alcohols, for example heptadecaethyleneoxycetanol, or condensation products of ethylene oxide with partial esters derived from fatty acids and a hexitol such as
25 polyoxyethylene sorbitol monooleate, or condensation products of ethylene oxide with partial esters derived from fatty acids and hexitol anhydrides, for example polyethylene sorbitan monooleate. The aqueous suspensions can also contain one or more preservatives, for example ethyl, or n-propyl p-hydroxybenzoate, one or more coloring agents, one or more flavoring agents, and one or more sweetening agents, such as
30 sucrose or saccharin.

Oily suspensions can be formulated by suspending the active ingredients in a vegetable oil, for example arachis oil, olive oil, sesame oil or coconut oil, or in a mineral oil such as liquid paraffin. The oily suspensions can contain a thickening agent, for example beeswax, hard paraffin or cetyl alcohol. Sweetening agents and flavoring agents can be added to provide palatable oral preparations. These compositions can be preserved by the addition of an anti-oxidant such as ascorbic acid

Dispersible powders and granules suitable for preparation of an aqueous suspension by the addition of water provide the active ingredient in admixture with a dispersing or wetting agent, suspending agent and one or more preservatives. Suitable dispersing or wetting agents or suspending agents are exemplified by those already mentioned above. Additional excipients, for example sweetening, flavoring and coloring agents, can also be present.

Pharmaceutical compositions of the invention can also be in the form of oil-in-water emulsions. The oily phase can be a vegetable oil or a mineral oil or mixtures of these. Suitable emulsifying agents can be naturally-occurring gums, for example gum acacia or gum tragacanth, naturally-occurring phosphatides, for example soy bean, lecithin, and esters or partial esters derived from fatty acids and hexitol, anhydrides, for example sorbitan monooleate, and condensation products of the said partial esters with ethylene oxide, for example polyoxyethylene sorbitan monooleate. The emulsions can also contain sweetening and flavoring agents.

Syrups and elixirs can be formulated with sweetening agents, for example glycerol, propylene glycol, sorbitol, glucose or sucrose. Such formulations can also contain a demulcent, a preservative and flavoring and coloring agents. The pharmaceutical compositions can be in the form of a sterile injectable aqueous or oleaginous suspension. This suspension can be formulated according to the known art using those suitable dispersing or wetting agents and suspending agents that have been mentioned above. The sterile injectable preparation can also be a sterile injectable solution or suspension in a non-toxic parentally acceptable diluent or solvent, for example as a solution in 1,3-butanediol. Among the acceptable vehicles and solvents that can be employed are water, Ringer's solution and isotonic sodium chloride solution. In addition, sterile, fixed oils are conventionally employed as a solvent or suspending medium. For this purpose, any

bland fixed oil can be employed including synthetic mono-or diglycerides. In addition, fatty acids such as oleic acid find use in the preparation of injectables.

The nucleic acid molecules of the invention can also be administered in the form of suppositories, *e.g.*, for rectal administration of the drug. These compositions can be prepared by mixing the drug with a suitable non-irritating excipient that is solid at ordinary temperatures but liquid at the rectal temperature and will therefore melt in the rectum to release the drug. Such materials include cocoa butter and polyethylene glycols.

Nucleic acid molecules of the invention can be administered parenterally in a sterile medium. The drug, depending on the vehicle and concentration used, can either be suspended or dissolved in the vehicle. Advantageously, adjuvants such as local anesthetics, preservatives and buffering agents can be dissolved in the vehicle.

Dosage levels of the order of from about 0.1 mg to about 140 mg per kilogram of body weight per day are useful in the treatment of the above-indicated conditions (about 0.5 mg to about 7 g per subject per day). The amount of active ingredient that can be combined with the carrier materials to produce a single dosage form varies depending upon the host treated and the particular mode of administration. Dosage unit forms generally contain between from about 1 mg to about 500 mg of an active ingredient.

It is understood that the specific dose level for any particular subject depends upon a variety of factors including the activity of the specific compound employed, the age, body weight, general health, sex, diet, time of administration, route of administration, and rate of excretion, drug combination and the severity of the particular disease undergoing therapy.

For administration to non-human animals, the composition can also be added to the animal feed or drinking water. It can be convenient to formulate the animal feed and drinking water compositions so that the animal takes in a therapeutically appropriate quantity of the composition along with its diet. It can also be convenient to present the composition as a premix for addition to the feed or drinking water.

The nucleic acid molecules of the present invention can also be administered to a subject in combination with other therapeutic compounds to increase the overall therapeutic effect. The use of multiple compounds to treat an indication can increase the beneficial effects while reducing the presence of side effects.

5 In one embodiment, the invention comprises compositions suitable for administering nucleic acid molecules of the invention to specific cell types. For example, the asialoglycoprotein receptor (ASGPr) (Wu and Wu, 1987, *J. Biol. Chem.* 262, 4429-4432) is unique to hepatocytes and binds branched galactose-terminal glycoproteins, such as asialoorosomucoid (ASOR). In another example, the folate
10 receptor is overexpressed in many cancer cells. Binding of such glycoproteins, synthetic glycoconjugates, or folates to the receptor takes place with an affinity that strongly depends on the degree of branching of the oligosaccharide chain, for example, triantennary structures are bound with greater affinity than biantennary or monoantennary chains (Baenziger and Fiete, 1980, *Cell*, 22, 611-620; Connolly *et al.*, 1982, *J. Biol.*
15 *Chem.*, 257, 939-945). Lee and Lee, 1987, *Glycoconjugate J.*, 4, 317-328, obtained this high specificity through the use of N-acetyl-D-galactosamine as the carbohydrate moiety, which has higher affinity for the receptor, compared to galactose. This "clustering effect" has also been described for the binding and uptake of mannosyl-terminating glycoproteins or glycoconjugates (Ponpipom *et al.*, 1981, *J. Med. Chem.*, 24, 1388-
20 1395). The use of galactose, galactosamine, or folate based conjugates to transport exogenous compounds across cell membranes can provide a targeted delivery approach to, for example, the treatment of liver disease, cancers of the liver, or other cancers. The use of bioconjugates can also provide a reduction in the required dose of therapeutic compounds required for treatment. Furthermore, therapeutic bioavailability,
25 pharmacodynamics, and pharmacokinetic parameters can be modulated through the use of nucleic acid bioconjugates of the invention. Non-limiting examples of such bioconjugates are described in Vargeese *et al.*, USSN 10/201,394, filed August 13, 2001; and Matulic-Adamic *et al.*, USSN 60/362,016, filed March 6, 2002.

Alternatively, certain siNA molecules of the instant invention can be expressed
30 within cells from eukaryotic promoters (e.g., Izant and Weintraub, 1985, *Science*, 229, 345; McGarry and Lindquist, 1986, *Proc. Natl. Acad. Sci.*, USA 83, 399; Scanlon *et al.*,

- 1991, *Proc. Natl. Acad. Sci. USA*, 88, 10591-5; Kashani-Sabet *et al.*, 1992, *Antisense Res. Dev.*, 2, 3-15; Dropulic *et al.*, 1992, *J. Virol.*, 66, 1432-41; Weerasinghe *et al.*, 1991, *J. Virol.*, 65, 5531-4; Ojwang *et al.*, 1992, *Proc. Natl. Acad. Sci. USA*, 89, 10802-6; Chen *et al.*, 1992, *Nucleic Acids Res.*, 20, 4581-9; Sarver *et al.*, 1990 *Science*, 247, 1222-1225; Thompson *et al.*, 1995, *Nucleic Acids Res.*, 23, 2259; Good *et al.*, 1997, *Gene Therapy*, 4, 45. Those skilled in the art realize that any nucleic acid can be expressed in eukaryotic cells from the appropriate DNA/RNA vector. The activity of such nucleic acids can be augmented by their release from the primary transcript by a enzymatic nucleic acid (Draper *et al.*, PCT WO 93/23569, and Sullivan *et al.*, PCT WO 94/02595; Ohkawa *et al.*, 1992, *Nucleic Acids Symp. Ser.*, 27, 15-6; Taira *et al.*, 1991, *Nucleic Acids Res.*, 19, 5125-30; Ventura *et al.*, 1993, *Nucleic Acids Res.*, 21, 3249-55; Chowrira *et al.*, 1994, *J. Biol. Chem.*, 269, 25856.

- In another aspect of the invention, RNA molecules of the present invention can be expressed from transcription units (see for example Couture *et al.*, 1996, *TIG.*, 12, 510) inserted into DNA or RNA vectors. The recombinant vectors can be DNA plasmids or viral vectors. siNA expressing viral vectors can be constructed based on, but not limited to, adeno-associated virus, retrovirus, adenovirus, or alphavirus. In another embodiment, pol III based constructs are used to express nucleic acid molecules of the invention (see for example Thompson, U.S. Pats. Nos. 5,902,880 and 6,146,886). The recombinant vectors capable of expressing the siNA molecules can be delivered as described above, and persist in target cells. Alternatively, viral vectors can be used that provide for transient expression of nucleic acid molecules. Such vectors can be repeatedly administered as necessary. Once expressed, the siNA molecule interacts with the target mRNA and generates an RNAi response. Delivery of siNA molecule expressing vectors can be systemic, such as by intravenous or intra-muscular administration, by administration to target cells ex-planted from a subject followed by reintroduction into the subject, or by any other means that would allow for introduction into the desired target cell (for a review see Couture *et al.*, 1996, *TIG.*, 12, 510).

- In one aspect the invention features an expression vector comprising a nucleic acid sequence encoding at least one siNA molecule of the instant invention. The expression vector can encode one or both strands of a siNA duplex, or a single self-complementary

strand that self hybridizes into a siNA duplex. The nucleic acid sequences encoding the siNA molecules of the instant invention can be operably linked in a manner that allows expression of the siNA molecule (see for example Paul *et al.*, 2002, *Nature Biotechnology*, 19, 505; Miyagishi and Taira, 2002, *Nature Biotechnology*, 19, 497; Lee *et al.*, 2002, *Nature Biotechnology*, 19, 500; and Novina *et al.*, 2002, *Nature Medicine*, advance online publication doi:10.1038/nm725).

In another aspect, the invention features an expression vector comprising: a) a transcription initiation region (*e.g.*, eukaryotic pol I, II or III initiation region); b) a transcription termination region (*e.g.*, eukaryotic pol I, II or III termination region); and
 10 c) a nucleic acid sequence encoding at least one of the siNA molecules of the instant invention, wherein said sequence is operably linked to said initiation region and said termination region in a manner that allows expression and/or delivery of the siNA molecule. The vector can optionally include an open reading frame (ORF) for a protein operably linked on the 5' side or the 3'-side of the sequence encoding the siNA of the
 15 invention; and/or an intron (intervening sequences).

Transcription of the siNA molecule sequences can be driven from a promoter for eukaryotic RNA polymerase I (pol I), RNA polymerase II (pol II), or RNA polymerase III (pol III). Transcripts from pol II or pol III promoters are expressed at high levels in all cells; the levels of a given pol II promoter in a given cell type depends on the nature
 20 of the gene regulatory sequences (enhancers, silencers, etc.) present nearby. Prokaryotic RNA polymerase promoters are also used, providing that the prokaryotic RNA polymerase enzyme is expressed in the appropriate cells (Elroy-Stein and Moss, 1990, *Proc. Natl. Acad. Sci. U S A*, 87, 6743-7; Gao and Huang 1993, *Nucleic Acids Res.*, 21, 2867-72; Lieber *et al.*, 1993, *Methods Enzymol.*, 217, 47-66; Zhou *et al.*, 1990, *Mol.*
 25 *Cell. Biol.*, 10, 4529-37). Several investigators have demonstrated that nucleic acid molecules expressed from such promoters can function in mammalian cells (*e.g.* Kashani-Sabet *et al.*, 1992, *Antisense Res. Dev.*, 2, 3-15; Ojwang *et al.*, 1992, *Proc. Natl. Acad. Sci. U S A*, 89, 10802-6; Chen *et al.*, 1992, *Nucleic Acids Res.*, 20, 4581-9; Yu *et al.*, 1993, *Proc. Natl. Acad. Sci. U S A*, 90, 6340-4; L'Huillier *et al.*, 1992, *EMBO*
 30 *J.*, 11, 4411-8; Lisiewicz *et al.*, 1993, *Proc. Natl. Acad. Sci. U. S. A*, 90, 8000-4; Thompson *et al.*, 1995, *Nucleic Acids Res.*, 23, 2259; Sullenger & Cech, 1993, *Science*,

262, 1566). More specifically, transcription units such as the ones derived from genes encoding U6 small nuclear (snRNA), transfer RNA (tRNA) and adenovirus VA RNA are useful in generating high concentrations of desired RNA molecules such as siNA in cells (Thompson *et al.*, *supra*; Couture and Stinchcomb, 1996, *supra*; Noonberg *et al.*, 1994, 5 *Nucleic Acid Res.*, 22, 2830; Noonberg *et al.*, U.S. Pat. No. 5,624,803; Good *et al.*, 1997, *Gene Ther.*, 4, 45; Beigelman *et al.*, International PCT Publication No. WO 96/18736. The above siNA transcription units can be incorporated into a variety of vectors for introduction into mammalian cells, including but not restricted to, plasmid DNA vectors, viral DNA vectors (such as adenovirus or adeno-associated virus vectors), or viral RNA 10 vectors (such as retroviral or alphavirus vectors) (for a review see Couture and Stinchcomb, 1996, *supra*).

In another aspect the invention features an expression vector comprising a nucleic acid sequence encoding at least one of the siNA molecules of the invention in a manner that allows expression of that siNA molecule. The expression vector comprises in one 15 embodiment; a) a transcription initiation region; b) a transcription termination region; and c) a nucleic acid sequence encoding at least one strand of the siNA molecule, wherein the sequence is operably linked to the initiation region and the termination region in a manner that allows expression and/or delivery of the siNA molecule.

In another embodiment the expression vector comprises: a) a transcription 20 initiation region; b) a transcription termination region; c) an open reading frame; and d) a nucleic acid sequence encoding at least one strand of a siNA molecule, wherein the sequence is operably linked to the 3'-end of the open reading frame and wherein the sequence is operably linked to the initiation region, the open reading frame and the termination region in a manner that allows expression and/or delivery of the siNA 25 molecule. In yet another embodiment, the expression vector comprises: a) a transcription initiation region; b) a transcription termination region; c) an intron; and d) a nucleic acid sequence encoding at least one siNA molecule, wherein the sequence is operably linked to the initiation region, the intron and the termination region in a manner which allows expression and/or delivery of the nucleic acid molecule.

30 In another embodiment, the expression vector comprises: a) a transcription initiation region; b) a transcription termination region; c) an intron; d) an open reading

- frame; and e) a nucleic acid sequence encoding at least one strand of a siNA molecule, wherein the sequence is operably linked to the 3'-end of the open reading frame and wherein the sequence is operably linked to the initiation region, the intron, the open reading frame and the termination region in a manner which allows expression and/or
- 5 delivery of the siNA molecule.

Expressed pseudogene target biology and biochemistry

- Pseudogenes have been considered as nonfunctional sequences of genomic DNA originally derived from functional genes. The assumption follows that all pseudogene mutations are selectively neutral and have equal probability to become fixed in the
- 10 population. However, pseudogenes that have been suitably investigated often exhibit functional roles, such as gene expression, gene regulation, generation of genetic (e.g., antibody, antigenic, and other) diversity (see for example Balakirev *et al.*, 2003, *Annu. Rev. Genet.*, 37, 123-51). A pseudogene is an evolutionary conserved gene copy that does not produce a functional, full-length protein. The human genome is estimated to
- 15 contain upwards of 20,000 pseudogenes. Although much effort has been devoted to understanding the function of such pseudogenes, their biological roles remain largely unknown. Some pseudogenes that are expressed have been associated with disease or developmental conditions. For example, Hirotsune *et al.*, 2003, *Nature*, 423, 91-96, report the role of an expressed pseudogene, specifically regulation of messenger-RNA
- 20 stability, in a transgene-insertion mouse mutant exhibiting polycystic kidneys and bone deformity. The transgene was integrated into the vicinity of the expressing pseudogene of Makorin1, referred to as Makorin1-p1. This insertion attenuated transcription of Makorin1-p1, resulting in destabilization of Makorin1 mRNA in trans by way of a cis-acting RNA decay element within the 5' region of Makorin1 that is homologous between
- 25 Makorin1 and Makorin1-p1. Either Makorin1 or Makorin1-p1 transgenes could rescue the expressed pseudogene phenotypes. These findings demonstrate a specific regulatory role of an expressed pseudogene, and point to the functional significance of non-coding RNAs (Hirotsune *et al.*, 2003, *Nature*, 423, 91-96). In a subsequent study, it was determined that 2-3% of human processed pseudogenes are expressed (Yano *et al.*, 2004,
- 30 *J. Mol. Med.*, 82(7), 414-22). Other reports of expressed pseudogenes are described in, for example Kandouz *et al.*, 2004, *Oncogene*, 23, 4763-70 (Connexin43); Yoshida *et al.*,

2003, *Hum Cell*, 16, 65-72 (Makorin1); and Perez Jurado *et al.*, 1998, *Hum Mol Genet*, 7, 325-34).

Examples:

The following are non-limiting examples showing the selection, isolation,
5 synthesis and activity of nucleic acids of the instant invention.

Example 1: Tandem synthesis of siNA constructs

Exemplary siNA molecules of the invention are synthesized in tandem using a
cleavable linker, for example, a succinyl-based linker. Tandem synthesis as described
herein is followed by a one-step purification process that provides RNAi molecules in
10 high yield. This approach is highly amenable to siNA synthesis in support of high
throughput RNAi screening, and can be readily adapted to multi-column or multi-well
synthesis platforms.

After completing a tandem synthesis of a siNA oligo and its complement in which
the 5'-terminal dimethoxytrityl (5'-O-DMT) group remains intact (trityl on synthesis), the
15 oligonucleotides are deprotected as described above. Following deprotection, the siNA
sequence strands are allowed to spontaneously hybridize. This hybridization yields a
duplex in which one strand has retained the 5'-O-DMT group while the complementary
strand comprises a terminal 5'-hydroxyl. The newly formed duplex behaves as a single
molecule during routine solid-phase extraction purification (Trityl-On purification) even
20 though only one molecule has a dimethoxytrityl group. Because the strands form a
stable duplex, this dimethoxytrityl group (or an equivalent group, such as other trityl
groups or other hydrophobic moieties) is all that is required to purify the pair of oligos,
for example, by using a C18 cartridge.

Standard phosphoramidite synthesis chemistry is used up to the point of
25 introducing a tandem linker, such as an inverted deoxy abasic succinate or glyceryl
succinate linker (see **Figure 1**) or an equivalent cleavable linker. A non-limiting
example of linker coupling conditions that can be used includes a hindered base such as
diisopropylethylamine (DIPA) and/or DMAP in the presence of an activator reagent such
as Bromotripyrrolidinophosphoniumhexafluorophosphate (PyBrOP). After the linker is

coupled, standard synthesis chemistry is utilized to complete synthesis of the second sequence leaving the terminal the 5'-O-DMT intact. Following synthesis, the resulting oligonucleotide is deprotected according to the procedures described herein and quenched with a suitable buffer, for example with 50mM NaOAc or 1.5M $\text{NH}_4\text{H}_2\text{CO}_3$.

- 5 Purification of the siNA duplex can be readily accomplished using solid phase extraction, for example, using a Waters C18 SepPak 1g cartridge conditioned with 1 column volume (CV) of acetonitrile, 2 CV H_2O , and 2 CV 50mM NaOAc. The sample is loaded and then washed with 1 CV H_2O or 50mM NaOAc. Failure sequences are eluted with 1 CV 14% ACN (Aqueous with 50mM NaOAc and 50mM NaCl). The
- 10 column is then washed, for example with 1 CV H_2O followed by on-column detritylation, for example by passing 1 CV of 1% aqueous trifluoroacetic acid (TFA) over the column, then adding a second CV of 1% aqueous TFA to the column and allowing to stand for approximately 10 minutes. The remaining TFA solution is removed and the column washed with H_2O followed by 1 CV 1M NaCl and additional
- 15 H_2O . The siNA duplex product is then eluted, for example, using 1 CV 20% aqueous CAN.

- Figure 2 provides an example of MALDI-TOF mass spectrometry analysis of a purified siNA construct in which each peak corresponds to the calculated mass of an individual siNA strand of the siNA duplex. The same purified siNA provides three
- 20 peaks when analyzed by capillary gel electrophoresis (CGE), one peak presumably corresponding to the duplex siNA, and two peaks presumably corresponding to the separate siNA sequence strands. Ion exchange HPLC analysis of the same siNA contract only shows a single peak. Testing of the purified siNA construct using a luciferase reporter assay described below demonstrated the same RNAi activity compared to siNA
 - 25 constructs generated from separately synthesized oligonucleotide sequence strands.

Example 2: Identification of potential siNA target sites in any RNA sequence

- The sequence of an RNA target of interest, such as a viral or human mRNA transcript, is screened for target sites, for example by using a computer folding algorithm. In a non-limiting example, the sequence of a gene or RNA gene transcript
- 30 derived from a database, such as Genbank, is used to generate siNA targets having

complementarity to the target. Such sequences can be obtained from a database, or can be determined experimentally as known in the art. Target sites that are known, for example, those target sites determined to be effective target sites based on studies with other nucleic acid molecules, for example ribozymes or antisense, or those targets known to be associated with a disease or condition such as those sites containing mutations or deletions, can be used to design siNA molecules targeting those sites. Various parameters can be used to determine which sites are the most suitable target sites within the target RNA sequence. These parameters include but are not limited to secondary or tertiary RNA structure, the nucleotide base composition of the target sequence, the degree of homology between various regions of the target sequence, or the relative position of the target sequence within the RNA transcript. Based on these determinations, any number of target sites within the RNA transcript can be chosen to screen siNA molecules for efficacy, for example by using *in vitro* RNA cleavage assays, cell culture, or animal models. In a non-limiting example, anywhere from 1 to 1000 target sites are chosen within the transcript based on the size of the siNA construct to be used. High throughput screening assays can be developed for screening siNA molecules using methods known in the art, such as with multi-well or multi-plate assays to determine efficient reduction in target gene expression.

Example 3: Selection of siNA molecule target sites in a RNA

- The following non-limiting steps can be used to carry out the selection of siNAs targeting a given gene sequence or transcript.
1. The target sequence is parsed *in silico* into a list of all fragments or subsequences of a particular length, for example 23 nucleotide fragments, contained within the target sequence. This step is typically carried out using a custom Perl script, but commercial sequence analysis programs such as Oligo, MacVector, or the GCG Wisconsin Package can be employed as well.
 2. In some instances the siNAs correspond to more than one target sequence; such would be the case for example in targeting different transcripts of the same gene, targeting different transcripts of more than one gene, or for targeting both the human gene and an animal homolog. In this case, a subsequence list of a particular length is

- generated for each of the targets, and then the lists are compared to find matching sequences in each list. The subsequences are then ranked according to the number of target sequences that contain the given subsequence; the goal is to find subsequences that are present in most or all of the target sequences. Alternately, the ranking can identify subsequences that are unique to a target sequence, such as a mutant target sequence. Such an approach would enable the use of siNA to target specifically the mutant sequence and not effect the expression of the normal sequence.
3. In some instances the siNA subsequences are absent in one or more sequences while present in the desired target sequence; such would be the case if the siNA targets a gene with a paralogous family member that is to remain untargeted. As in case 2 above, a subsequence list of a particular length is generated for each of the targets, and then the lists are compared to find sequences that are present in the target gene but are absent in the untargeted paralog.
4. The ranked siNA subsequences can be further analyzed and ranked according to GC content. A preference can be given to sites containing 30-70% GC, with a further preference to sites containing 40-60% GC.
5. The ranked siNA subsequences can be further analyzed and ranked according to self-folding and internal hairpins. Weaker internal folds are preferred; strong hairpin structures are to be avoided.
6. The ranked siNA subsequences can be further analyzed and ranked according to whether they have runs of GGG or CCC in the sequence. GGG (or even more Gs) in either strand can make oligonucleotide synthesis problematic and can potentially interfere with RNAi activity, so it is avoided whenever better sequences are available. CCC is searched in the target strand because that will place GGG in the antisense strand.
7. The ranked siNA subsequences can be further analyzed and ranked according to whether they have the dinucleotide UU (uridine dinucleotide) on the 3'-end of the sequence, and/or AA on the 5'-end of the sequence (to yield 3' UU on the antisense sequence). These sequences allow one to design siNA molecules with terminal TT thymidine dinucleotides.

8. Four or five target sites are chosen from the ranked list of subsequences as described above. For example, in subsequences having 23 nucleotides, the right 21 nucleotides of each chosen 23-mer subsequence are then designed and synthesized for the upper (sense) strand of the siNA duplex, while the reverse complement of the left 21 nucleotides of each chosen 23-mer subsequence are then designed and synthesized for the lower (antisense) strand of the siNA duplex. If terminal TT residues are desired for the sequence (as described in paragraph 7), then the two 3' terminal nucleotides of both the sense and antisense strands are replaced by TT prior to synthesizing the oligos.
9. The siNA molecules are screened in an *in vitro*, cell culture or animal model system to identify the most active siNA molecule or the most preferred target site within the target RNA sequence.
10. Other design considerations can be used when selecting target nucleic acid sequences, see, for example, Reynolds *et al.*, 2004, *Nature Biotechnology Advanced Online Publication*, 1 February 2004, doi:10.1038/nbt936 and Ui-Tei *et al.*, 2004, *Nucleic Acids Research*, 32, doi:10.1093/nar/gkh247.

- In an alternate approach, a pool of siNA constructs specific to a target polynucleotide sequence is used to screen for target sites in cells expressing target RNA. The general strategy used in this approach is shown in Figure 9. Cells expressing target RNA are transfected with the pool of siNA constructs and cells that demonstrate a phenotype associated with target inhibition are sorted. The pool of siNA constructs can be expressed from transcription cassettes inserted into appropriate vectors (see for example Figure 7 and Figure 8). The siNA from cells demonstrating a positive phenotypic change (e.g., decreased proliferation, decreased RNA levels, decreased protein expression), are sequenced to determine the most suitable target site(s) within the target RNA sequence.

Example 4: Targeted siNA design

- siNA target sites were chosen by analyzing sequences of the target polynucleotide and optionally prioritizing the target sites on the basis of folding (structure of any given sequence analyzed to determine siNA accessibility to the target), by using a library of

siNA molecules as described in Example 3, or alternately by using an *in vitro* siNA system as described in Example 6 herein. siNA molecules are designed that could bind each target and are optionally individually analyzed by computer folding to assess whether the siNA molecule can interact with the target sequence. Varying the length of the siNA molecules can be chosen to optimize activity. Generally, a sufficient number of complementary nucleotide bases are chosen to bind to, or otherwise interact with, the target RNA, but the degree of complementarity can be modulated to accommodate siNA duplexes or varying length or base composition. By using such methodologies, siNA molecules can be designed to target sites within any known RNA sequence, for example those RNA sequences corresponding to the any gene transcript.

Chemically modified siNA constructs are designed to provide nuclease stability for systemic administration *in vivo* and/or improved pharmacokinetic, localization, and delivery properties while preserving the ability to mediate RNAi activity. Chemical modifications as described herein are introduced synthetically using synthetic methods described herein and those generally known in the art. The synthetic siNA constructs are then assayed for nuclease stability in serum and/or cellular/tissue extracts (e.g. liver extracts). The synthetic siNA constructs are also tested in parallel for RNAi activity using an appropriate assay, such as a luciferase reporter assay as described herein or another suitable assay that can quantify RNAi activity. Synthetic siNA constructs that possess both nuclease stability and RNAi activity can be further modified and re-evaluated in stability and activity assays. The chemical modifications of the stabilized active siNA constructs can then be applied to any siNA sequence targeting any chosen RNA and used, for example, in target screening assays to pick lead siNA compounds for therapeutic development (see for example **Figure 11**).

25 Example 5: Chemical Synthesis and Purification of siNA

siNA molecules can be designed to interact with various sites in the RNA message, for example, target sequences within the RNA sequences described herein. The sequence of one strand of the siNA molecule(s) is complementary to the target site sequences described above. The siNA molecules can be chemically synthesized using methods described herein. Inactive siNA molecules that are used as control sequences can be synthesized by scrambling the sequence of the siNA molecules such that it is not

complementary to the target sequence. Generally, siNA constructs can be synthesized using solid phase oligonucleotide synthesis methods as described herein (see for example Usman *et al.*, US Patent Nos. 5,804,683; 5,831,071; 5,998,203; 6,117,657; 6,353,098; 6,362,323; 6,437,117; 6,469,158; Scaringe *et al.*, US Patent Nos. 6,111,086; 6,008,400; 6,111,086 all incorporated by reference herein in their entirety).

In a non-limiting example, RNA oligonucleotides are synthesized in a stepwise fashion using the phosphoramidite chemistry as is known in the art. Standard phosphoramidite chemistry involves the use of nucleosides comprising any of 5'-O-dimethoxytrityl, 2'-O-tert-butylidimethylsilyl, 3'-O-2-Cyanoethyl N,N-diisopropylphosphoroamidite groups, and exocyclic amine protecting groups (e.g. N6-benzoyl adenosine, N4 acetyl cytidine, and N2-isobutryl guanosine). Alternately, 2'-O-Silyl Ethers can be used in conjunction with acid-labile 2'-O-orthoester protecting groups in the synthesis of RNA as described by Scaringe *supra*. Differing 2' chemistries can require different protecting groups, for example 2'-deoxy-2'-amino nucleosides can utilize N-phthaloyl protection as described by Usman *et al.*, US Patent 5,631,360, incorporated by reference herein in its entirety).

During solid phase synthesis, each nucleotide is added sequentially (3'- to 5'-direction) to the solid support-bound oligonucleotide. The first nucleoside at the 3'-end of the chain is covalently attached to a solid support (e.g., controlled pore glass or polystyrene) using various linkers. The nucleotide precursor, a ribonucleoside phosphoramidite, and activator are combined resulting in the coupling of the second nucleoside phosphoramidite onto the 5'-end of the first nucleoside. The support is then washed and any unreacted 5'-hydroxyl groups are capped with a capping reagent such as acetic anhydride to yield inactive 5'-acetyl moieties. The trivalent phosphorus linkage is then oxidized to a more stable phosphate linkage. At the end of the nucleotide addition cycle, the 5'-O-protecting group is cleaved under suitable conditions (e.g., acidic conditions for trityl-based groups and Fluoride for silyl-based groups). The cycle is repeated for each subsequent nucleotide.

Modification of synthesis conditions can be used to optimize coupling efficiency, for example by using differing coupling times, differing reagent/phosphoramidite concentrations, differing contact times, differing solid supports and solid support linker

chemistries depending on the particular chemical composition of the siNA to be synthesized. Deprotection and purification of the siNA can be performed as is generally described in Usman *et al.*, US 5,831,071, US 6,353,098, US 6,437,117, and Bellon *et al.*, US 6,054,576, US 6,162,909, US 6,303,773, or Scaringe *supra*, incorporated by
5 reference herein in their entireties. Additionally, deprotection conditions can be modified to provide the best possible yield and purity of siNA constructs. For example, applicant has observed that oligonucleotides comprising 2'-deoxy-2'-fluoro nucleotides can degrade under inappropriate deprotection conditions. Such oligonucleotides are deprotected using aqueous methylamine at about 35°C for 30 minutes. If the 2'-deoxy-
10 2'-fluoro containing oligonucleotide also comprises ribonucleotides, after deprotection with aqueous methylamine at about 35°C for 30 minutes, TEA-HF is added and the reaction maintained at about 65°C for an additional 15 minutes.

Example 6: RNAi *in vitro* assay to assess siNA activity

An *in vitro* assay that recapitulates RNAi in a cell-free system is used to evaluate
15 siNA constructs targeting target RNA. The assay comprises the system described by Tuschl *et al.*, 1999, *Genes and Development*, 13, 3191-3197 and Zamore *et al.*, 2000, *Cell*, 101, 25-33 adapted for use with target RNA. A *Drosophila* extract derived from syncytial blastoderm is used to reconstitute RNAi activity *in vitro*. Target RNA is generated via *in vitro* transcription from an appropriate target expressing plasmid using
20 T7 RNA polymerase or via chemical synthesis as described herein. Sense and antisense siNA strands (for example 20 uM each) are annealed by incubation in buffer (such as 100 mM potassium acetate, 30 mM HEPES-KOH, pH 7.4, 2 mM magnesium acetate) for 1 minute at 90°C followed by 1 hour at 37°C, then diluted in lysis buffer (for example 100 mM potassium acetate, 30 mM HEPES-KOH at pH 7.4, 2mM magnesium acetate).
25 Annealing can be monitored by gel electrophoresis on an agarose gel in TBE buffer and stained with ethidium bromide. The *Drosophila* lysate is prepared using zero to two-hour-old embryos from Oregon R flies collected on yeasted molasses agar that are dechorionated and lysed. The lysate is centrifuged and the supernatant isolated. The assay comprises a reaction mixture containing 50% lysate [vol/vol], RNA (10-50 pM
30 final concentration), and 10% [vol/vol] lysis buffer containing siNA (10 nM final concentration). The reaction mixture also contains 10 mM creatine phosphate, 10 ug/ml

creatine phosphokinase, 100 μ M GTP, 100 μ M UTP, 100 μ M CTP, 500 μ M ATP, 5 mM DTT, 0.1 U/ μ L RNasin (Promega), and 100 μ M of each amino acid. The final concentration of potassium acetate is adjusted to 100 mM. The reactions are pre-assembled on ice and preincubated at 25° C for 10 minutes before adding RNA, then

5 incubated at 25° C for an additional 60 minutes. Reactions are quenched with 4 volumes of 1.25 x Passive Lysis Buffer (Promega). Target RNA cleavage is assayed by RT-PCR analysis or other methods known in the art and are compared to control reactions in which siNA is omitted from the reaction.

Alternately, internally-labeled target RNA for the assay is prepared by *in vitro*

10 transcription in the presence of [α -³²P] CTP, passed over a G50 Sephadex column by spin chromatography and used as target RNA without further purification. Optionally, target RNA is 5'-³²P-end labeled using T4 polynucleotide kinase enzyme. Assays are performed as described above and target RNA and the specific RNA cleavage products generated by RNAi are visualized on an autoradiograph of a gel. The percentage of

15 cleavage is determined by PHOSPHOR IMAGER® (autoradiography) quantitation of bands representing intact control RNA or RNA from control reactions without siNA and the cleavage products generated by the assay.

In one embodiment, this assay is used to determine target sites in the target RNA target for siNA mediated RNAi cleavage, wherein a plurality of siNA constructs are

20 screened for RNAi mediated cleavage of the target RNA, for example, by analyzing the assay reaction by electrophoresis of labeled target RNA, or by northern blotting, as well as by other methodology well known in the art.

Example 7: Nucleic acid inhibition of target RNA

siNA molecules targeted to the human target RNA are designed and synthesized as

25 described above. These nucleic acid molecules can be tested for cleavage activity *in vivo*, for example, using the following procedure.

Two formats are used to test the efficacy of siNAs of the invention. First, the reagents are tested in cell culture to determine the extent of RNA and protein inhibition. siNA reagents are selected against the target as described herein. RNA inhibition is

- measured after delivery of these reagents by a suitable transfection agent to cells. Relative amounts of target RNA are measured versus actin using real-time PCR monitoring of amplification (eg., ABI 7700 TAQMAN® (real-time PCR monitoring of amplification)). A comparison is made to a mixture of oligonucleotide sequences
- 5 made to unrelated targets or to a randomized siNA control with the same overall length and chemistry, but randomly substituted at each position. Primary and secondary lead reagents are chosen for the target and optimization performed. After an optimal transfection agent concentration is chosen, a RNA time-course of inhibition is performed with the lead siNA molecule. In addition, a cell-plating format can be used to determine
- 10 RNA inhibition.

Delivery of siNA to Cells

- Cells (e.g., HEK293/HEK293T, HeLa, A549, A375 cells) are seeded, for example, at 1×10^5 cells per well of a six-well dish in EGM-2 (BioWhittaker) the day before transfection. siNA (final concentration, for example 20nM) and cationic lipid (e.g., final
- 15 concentration $2 \mu\text{g/ml}$) are complexed in EGM basal media (Bio Whittaker) at 37°C for 30 minutes in polystyrene tubes. Following vortexing, the complexed siNA is added to each well and incubated for the times indicated. For initial optimization experiments, cells are seeded, for example, at 1×10^3 in 96 well plates and siNA complex added as described. Efficiency of delivery of siNA to cells is determined using a fluorescent siNA
- 20 complexed with lipid. Cells in 6-well dishes are incubated with siNA for 24 hours, rinsed with PBS and fixed in 2% paraformaldehyde for 15 minutes at room temperature. Uptake of siNA is visualized using a fluorescent microscope.

TAQMAN® (real-time PCR monitoring of amplification) and Lightcycler quantification of mRNA

- 25 Total RNA is prepared from cells following siNA delivery, for example, using Qiagen RNA purification kits for 6-well or Rneasy extraction kits for 96-well assays. For TAQMAN® analysis (real-time PCR monitoring of amplification), dual-labeled probes are synthesized with the reporter dye, FAM or JOE, covalently linked at the 5'-end and the quencher dye TAMRA conjugated to the 3'-end. One-step RT-PCR amplifications
- 30 are performed on, for example, an ABI PRISM 7700 Sequence Detector using $50 \mu\text{l}$

reactions consisting of 10 μ l total RNA, 100 nM forward primer, 900 nM reverse primer, 100 nM probe, 1X TaqMan PCR reaction buffer (PE-Applied Biosystems), 5.5 mM $MgCl_2$, 300 μ M each dATP, dCTP, dGTP, and dTTP, 10U RNase Inhibitor (Promega), 1.25U AMPLITAQ GOLD® (DNA polymerase) (PE-Applied Biosystems) and 10U M-MLV Reverse Transcriptase (Promega). The thermal cycling conditions can consist of 30 minutes at 48°C, 10 minutes at 95°C, followed by 40 cycles of 15 seconds at 95°C and 1 minute at 60°C. Quantitation of mRNA levels is determined relative to standards generated from serially diluted total cellular RNA (300, 100, 33, 11 ng/rxn) and normalizing to β -actin or GAPDH mRNA in parallel TAQMAN® reactions (real-time PCR monitoring of amplification). For each gene of interest an upper and lower primer and a fluorescently labeled probe are designed. Real time incorporation of SYBR Green I dye into a specific PCR product can be measured in glass capillary tubes using a lightcycler. A standard curve is generated for each primer pair using control cRNA. Values are represented as relative expression to GAPDH in each sample.

15 Western blotting

Nuclear extracts can be prepared using a standard micro preparation technique (see for example Andrews and Faller, 1991, *Nucleic Acids Research*, 19, 2499). Protein extracts from supernatants are prepared, for example using TCA precipitation. An equal volume of 20% TCA is added to the cell supernatant, incubated on ice for 1 hour and pelleted by centrifugation for 5 minutes. Pellets are washed in acetone, dried and resuspended in water. Cellular protein extracts are run on a 10% Bis-Tris NuPage (nuclear extracts) or 4-12% Tris-Glycine (supernatant extracts) polyacrylamide gel and transferred onto nitro-cellulose membranes. Non-specific binding can be blocked by incubation, for example, with 5% non-fat milk for 1 hour followed by primary antibody for 16 hour at 4°C. Following washes, the secondary antibody is applied, for example (1:10,000 dilution) for 1 hour at room temperature and the signal detected with SuperSignal reagent (Pierce).

Example 8: Models useful to evaluate the down-regulation of gene expression

Evaluating the efficacy of siNA molecules of the invention in animal models is an important prerequisite to human clinical trials. Various animal models of cancer,

proliferative, inflammatory, autoimmune, neurologic, ocular, respiratory, metabolic, etc. diseases, conditions, or disorders as are known in the art can be adapted for use for pre-clinical evaluation of the efficacy of nucleic acid compositions of the invention in modulating target gene expression toward therapeutic, cosmetic, or research use.

5 Example 9: RNAi mediated inhibition of target gene expression

siNA constructs (are tested for efficacy in reducing target RNA expression in cells, (e.g., HEK293/HEK293A, HeLa, A549, A375 cells). Cells are plated approximately 24 hours before transfection in 96-well plates at 5,000-7,500 cells/well, 100 μ l/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs are mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 μ l/well and incubated for 20 minutes at room temperature. The siNA transfection mixtures are added to cells to give a final siNA concentration of 25 nM in a volume of 150 μ l. Each siNA transfection mixture is added to 3 wells for triplicate siNA treatments. Cells are incubated at 37° for 24 hours in the continued presence of the siNA transfection mixture. At 24 hours, RNA is prepared from each well of treated cells. The supernatants with the transfection mixtures are first removed and discarded, then the cells are lysed and RNA prepared from each well. Target gene expression following treatment is evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data is averaged and the standard deviations determined for each treatment. Normalized data are graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs is determined.

Example 10: Indications

Particular conditions and disease states that can be associated with gene expression modulation include, but are not limited to proliferative, inflammatory, autoimmune, neurologic, ocular, respiratory, metabolic etc. diseases, conditions, or disorders as described herein or otherwise known in the art, and any other diseases, conditions or disorders that are related to or will respond to the levels of a target (e.g., target protein or target polynucleotide) in a cell or tissue, alone or in combination with other therapies.

Those skilled in the art will recognize that other drugs such as anti-cancer compounds and therapies can be similarly be readily combined with the nucleic acid molecules of the instant invention (*e.g.* ribozymes and antisense molecules) and are hence within the scope of the instant invention. Such compounds and therapies are well known in the art. For combination therapy, the nucleic acids of the invention are prepared in one of two ways. First, the agents are physically combined in a preparation of nucleic acid and chemotherapeutic agent, such as a mixture of a nucleic acid of the invention encapsulated in liposomes and ifosfamide in a solution for intravenous administration, wherein both agents are present in a therapeutically effective concentration (*e.g.*, ifosfamide in solution to deliver 1000-1250 mg/m²/day and liposome-associated nucleic acid of the invention in the same solution to deliver 0.1-100 mg/kg/day). Alternatively, the agents are administered separately but simultaneously in their respective effective doses (*e.g.*, 1000-1250 mg/m²/d ifosfamide and 0.1 to 100 mg/kg/day nucleic acid of the invention).

15 Example 11: Diagnostic uses

The siRNA molecules of the invention can be used in a variety of diagnostic applications, such as in the identification of molecular targets (*e.g.*, RNA) in a variety of applications, for example, in clinical, industrial, environmental, agricultural and/or research settings. Such diagnostic use of siRNA molecules involves utilizing reconstituted RNAi systems, for example, using cellular lysates or partially purified cellular lysates. siRNA molecules of this invention can be used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of endogenous or exogenous, for example viral, RNA in a cell. The close relationship between siRNA activity and the structure of the target RNA allows the detection of mutations in any region of the molecule, which alters the base-pairing and three-dimensional structure of the target RNA. By using multiple siRNA molecules described in this invention, one can map nucleotide changes, which are important to RNA structure and function *in vitro*, as well as in cells and tissues. Cleavage of target RNAs with siRNA molecules can be used to inhibit gene expression and define the role of specified gene products in the progression of disease or infection. In this manner, other genetic targets can be defined as important mediators of the disease. These experiments will lead to better treatment of

- the disease progression by affording the possibility of combination therapies (e.g., multiple siNA molecules targeted to different genes, siNA molecules coupled with known small molecule inhibitors, or intermittent treatment with combinations siNA molecules and/or other chemical or biological molecules). Other *in vitro* uses of siNA
- 5 molecules of this invention are well known in the art, and include detection of the presence of mRNAs associated with a disease, infection, or related condition. Such RNA is detected by determining the presence of a cleavage product after treatment with a siNA using standard methodologies, for example, fluorescence resonance emission transfer (FRET).
- 10 In a specific example, siNA molecules that cleave only wild-type or mutant forms of the target RNA are used for the assay. The first siNA molecules (*i.e.*, those that cleave only wild-type forms of target RNA) are used to identify wild-type RNA present in the sample and the second siNA molecules (*i.e.*, those that cleave only mutant forms of target RNA) are used to identify mutant RNA in the sample. As reaction controls,
- 15 synthetic substrates of both wild-type and mutant RNA are cleaved by both siNA molecules to demonstrate the relative siNA efficiencies in the reactions and the absence of cleavage of the "non-targeted" RNA species. The cleavage products from the synthetic substrates also serve to generate size markers for the analysis of wild-type and mutant RNAs in the sample population. Thus, each analysis requires two siNA
- 20 molecules, two substrates and one unknown sample, which is combined into six reactions. The presence of cleavage products is determined using an RNase protection assay so that full-length and cleavage fragments of each RNA can be analyzed in one lane of a polyacrylamide gel. It is not absolutely required to quantify the results to gain insight into the expression of mutant RNAs and putative risk of the desired phenotypic
- 25 changes in target cells. The expression of mRNA whose protein product is implicated in the development of the phenotype (*i.e.*, disease related or infection related) is adequate to establish risk. If probes of comparable specific activity are used for both transcripts, then a qualitative comparison of RNA levels is adequate and decreases the cost of the initial diagnosis. Higher mutant form to wild-type ratios are correlated with higher risk
- 30 whether RNA levels are compared qualitatively or quantitatively.

All patents and publications mentioned in the specification are indicative of the levels of skill of those skilled in the art to which the invention pertains. All references cited in this disclosure are incorporated by reference to the same extent as if each reference had been incorporated by reference in its entirety individually.

5 One skilled in the art would readily appreciate that the present invention is well adapted to carry out the objects and obtain the ends and advantages mentioned, as well as those inherent therein. The methods and compositions described herein as presently representative of preferred embodiments are exemplary and are not intended as limitations on the scope of the invention. Changes therein and other uses will occur to
10 those skilled in the art, which are encompassed within the spirit of the invention, are defined by the scope of the claims.

It will be readily apparent to one skilled in the art that varying substitutions and modifications can be made to the invention disclosed herein without departing from the scope and spirit of the invention. Thus, such additional embodiments are within the
15 scope of the present invention and the following claims. The present invention teaches one skilled in the art to test various combinations and/or substitutions of chemical modifications described herein toward generating nucleic acid constructs with improved activity for mediating RNAi activity. Such improved activity can comprise improved stability, improved bioavailability, and/or improved activation of cellular responses
20 mediating RNAi. Therefore, the specific embodiments described herein are not limiting and one skilled in the art can readily appreciate that specific combinations of the modifications described herein can be tested without undue experimentation toward identifying siRNA molecules with improved RNAi activity.

The invention illustratively described herein suitably can be practiced in the
25 absence of any element or elements, limitation or limitations that are not specifically disclosed herein. Thus, for example, in each instance herein any of the terms "comprising", "consisting essentially of", and "consisting of" may be replaced with either of the other two terms. The terms and expressions which have been employed are used as terms of description and not of limitation, and there is no intention that in the use of
30 such terms and expressions of excluding any equivalents of the features shown and described or portions thereof, but it is recognized that various modifications are possible

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within the scope of the invention claimed. Thus, it should be understood that although the present invention has been specifically disclosed by preferred embodiments, optional features, modification and variation of the concepts herein disclosed may be resorted to by those skilled in the art, and that such modifications and variations are considered to be

5 within the scope of this invention as defined by the description and the appended claims.

In addition, where features or aspects of the invention are described in terms of Markush groups or other grouping of alternatives, those skilled in the art will recognize that the invention is also thereby described in terms of any individual member or subgroup of members of the Markush group or other group.

NM_000536 Homo sapiens recombination activating gene 2 (RAG2), mRNA
 NM_000801 Homo sapiens hepatocyte growth factor (hepatopoietin A; scatter factor) (HGF), mRNA
 NM_000940 Homo sapiens paraoxonase 3 (PON3), mRNA
 NM_000941 Homo sapiens P450 (cytochrome) oxidoreductase (POR), mRNA
 NM_000953 Homo sapiens prostaglandin D2 receptor (DP) (PTGDR), mRNA
 NM_0010011 Homo sapiens Intersectin 1 (SH3 domain protein) (ITSN1), transcript variant
 NM_00100111 Homo sapiens transient receptor potential cation channel, subfamily M, member
 NM_0010012 Homo sapiens solute carrier family 2 (facilitated glucose transporter), member
 NM_0010013 Homo sapiens trypsin X3 (TRY1), mRNA
 NM_0010013 Homo sapiens ATPase, Ca++ transporting, plasma membrane 1 (ATP2B1), transcript
 NM_0010013 Homo sapiens similar to ubiquitin-specific protease 17-like protein (LOC4014
 NM_0010013 Homo sapiens Kazal type serine protease inhibitor 5-like 2 (SPINKSL2), mRNA
 NM_0010013 Homo sapiens protein kinase C substrate 80K-H (PRKCSH), transcript variant
 NM_0010013 Homo sapiens chromosome 10 open reading frame 74 (C10orf74), mRNA
 NM_0010013 Homo sapiens ATPase, Ca++ transporting, plasma membrane 2 (ATP2B2), transcript
 NM_0010013 Homo sapiens cytochrome b5 reductase b5R-2 (CYB5R2), transcript variant
 NM_0010013 Homo sapiens biogenesis of lysosome-related organelles complex-1, subunit
 NM_0010013 Homo sapiens MGC27121 gene (MGC27121), mRNA
 NM_0010013 Homo sapiens ATPase, Ca++ transporting, plasma membrane 3 (ATP2B3), transcript
 NM_0010013 Homo sapiens claudin 20 (CLDN20), mRNA
 NM_0010013 Homo sapiens NFkB inhibitor interacting Ras-like 2 (NKIRAS2), transcript variant
 NM_0010013 Homo sapiens CD44 antigen (homing function and Indian blood group system)
 NM_0010013 Homo sapiens CD44 antigen (homing function and Indian blood group system)
 NM_0010013 Homo sapiens CD44 antigen (homing function and Indian blood group system)
 NM_0010013 Homo sapiens CD44 antigen (homing function and Indian blood group system)
 NM_0010013 Homo sapiens HCG3 gene (HCG3), mRNA
 NM_0010013 Homo sapiens LIM domain only 3 (rhombotin-like 2) (LMO3), transcript variant
 NM_0010013 Homo sapiens ATPase, Ca++ transporting, plasma membrane 4 (ATP2B4), transcript
 NM_0010014 Homo sapiens hypothetical protein MGC24381 (MGC24381), mRNA
 NM_0010014 Homo sapiens similar to Zinc finger protein 208 (LOC163223), mRNA
 NM_0010014 Homo sapiens family with sequence similarity 26, member C (FAM26C), mRNA
 NM_0010014 Homo sapiens similar to F-box only protein 2 (LOC342897), mRNA
 NM_0010014 Homo sapiens zinc finger protein 429 (ZNF429), mRNA
 NM_0010014 Homo sapiens similar to TBC1 domain family, member 3, centromeric (LOC4
 NM_0010014 Homo sapiens similar to TBC1 domain family, member 3, telomeric (MGC44
 NM_0010014 Homo sapiens SMAD, mothers against DPP homolog 5 (Drosophila) (SMAD1
 NM_0010014 Homo sapiens SMAD, mothers against DPP homolog 5 (Drosophila) (SMAD1
 NM_0010014 Homo sapiens troponin T2, cardiac (TNNT2), transcript variant 2, mRNA
 NM_0010014 Homo sapiens troponin T2, cardiac (TNNT2), transcript variant 3, mRNA
 NM_0010014 Homo sapiens troponin T2, cardiac (TNNT2), transcript variant 4, mRNA
 NM_0010014 Homo sapiens syntaxin 16 (STX16), transcript variant 1, mRNA
 NM_0010014 Homo sapiens syntaxin 16 (STX16), transcript variant 3, mRNA
 NM_0010014 Homo sapiens chemokine (C-C motif) ligand 4-like (CCL4L), mRNA
 NM_0010014 Homo sapiens similar to RIKEN cDNA 4921524J17 (LOC388272), mRNA
 NM_0010014 Homo sapiens chemokine (C-C motif) ligand 3-like, centromeric (MGC12815
 NM_0010014 Homo sapiens lanosterol synthase (2,3-oxidosqualene-lanosterol cyclase) (L
 NM_0010014 Homo sapiens solute carrier family 35, member E4 (SLC35E4), mRNA
 NM_0010014 Homo sapiens hypothetical protein FLJ11011 (FLJ11011), transcript variant
 NM_0010014 Homo sapiens hypothetical protein FLJ11011 (FLJ11011), transcript variant
 NM_0010014 Homo sapiens zinc finger, DHHC domain containing 13 (ZDHHC13), transcript
 NM_0010014 Homo sapiens phosphotriesterase related (PTER), transcript variant 1, mRNA
 NM_0010014 Homo sapiens ATPase, Ca++ transporting, type 2C, member 1 (ATP2C1), transcript
 NM_0010014 Homo sapiens ATPase, Ca++ transporting, type 2C, member 1 (ATP2C1), transcript
 NM_0010014 Homo sapiens ATPase, Ca++ transporting, type 2C, member 1 (ATP2C1), transcript
 NM_0010015 Homo sapiens synuclein, beta (SNCB), transcript variant 1, mRNA
 NM_0010015 Homo sapiens NADH dehydrogenase (ubiquinone) flavoprotein 3, 10kDa (ND
 NM_0010015 Homo sapiens hepatoma-derived growth factor-related protein 2 (HDGF2), transcript

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NM_0010015: Homo sapiens UDP-glucose pyrophosphorylase 2 (UGP2), transcript variant
 NM_0010015: Homo sapiens tagalin (TAGLN), transcript variant 1, mRNA
 NM_0010015: Homo sapiens RAR-related orphan receptor C (RORC), transcript variant 2, i
 NM_0010015: Homo sapiens transmembrane 6 superfamily member 2 (TM6SF2), transcript
 NM_0010015: Homo sapiens CD36 antigen (collagen type I receptor, thrombospondin rece)
 NM_0010015: Homo sapiens CD36 antigen (collagen type I receptor, thrombospondin rece)
 NM_0010015: Homo sapiens growth factor receptor-bound protein 10 (GRB10), transcript v
 NM_0010015: Homo sapiens growth factor receptor-bound protein 10 (GRB10), transcript v
 NM_0010015: Homo sapiens chromosome 9 open reading frame 103 (C9orf103), mRNA
 NM_0010015: Homo sapiens LEM domain containing 1 (LEMD1), mRNA
 NM_0010015: Homo sapiens germ and embryonic stem cell enriched protein STELLA (STE
 NM_0010015: Homo sapiens growth factor receptor-bound protein 10 (GRB10), transcript v
 NM_0010015: Homo sapiens galactokinase 2 (GALK2), transcript variant 2, mRNA
 NM_0010015: Homo sapiens similar to growth differentiation factor 16 (LOC392255), mRNA/
 NM_0010015: Homo sapiens golgi associated, gamma adaptin ear containing, ARF binding
 NM_0010015: Homo sapiens golgi associated, gamma adaptin ear containing, ARF binding
 NM_0010015: Homo sapiens translocase of inner mitochondrial membrane 50 homolog (ye
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 2, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 3, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 4, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 5, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 6, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 7, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 8, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 9, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 10, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 11, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 12, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 13, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 14, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 15, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 16, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 17, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 18, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 19, mRNA
 NM_0010015: Homo sapiens phosphodiesterase 9A (PDE9A), transcript variant 20, mRNA
 NM_0010015: Homo sapiens ATPase, Na⁺/K⁺ transporting, alpha 1 polypeptide (ATP1A1),
 NM_0010016: Homo sapiens mediator of RNA polymerase II transcription, subunit 8 homol
 NM_0010016: Homo sapiens mediator of RNA polymerase II transcription, subunit 8 homol
 NM_0010016: Homo sapiens mediator of RNA polymerase II transcription, subunit 8 homol
 NM_0010016: Homo sapiens similar to hypothetical protein 9530023G02 (MGC90512), mR
 NM_0010016: Homo sapiens olfactory receptor, family 9, subfamily A, member 4 (OR9A4),
 NM_0010016: Homo sapiens olfactory receptor, family 9, subfamily A, member 2 (OR9A2),
 NM_0010016: Homo sapiens olfactory receptor, family 2, subfamily A, member 14 (OR2A1
 NM_0010016: Homo sapiens hypothetical protein LOC144363 (LOC144363), mRNA
 NM_0010016: Homo sapiens hypothetical protein LOC155054 (LOC155054), mRNA
 NM_0010016: Homo sapiens FLJ16636 protein (FLJ16636), mRNA
 NM_0010016: Homo sapiens hypothetical LOC256349 (ba9F11.1), mRNA
 NM_0010016: Homo sapiens hypothetical protein LOC339745 (LOC339745), mRNA
 NM_0010016: Homo sapiens FLJ16008 protein (FLJ16008), mRNA
 NM_0010016: Homo sapiens prostate cancer associated protein 5 (PCANAP5), transcript v
 NM_0010016: Homo sapiens olfactory receptor, family 6, subfamily V, member 1 (OR6V1),
 NM_0010016: Homo sapiens FLJ26175 protein (FLJ26175), mRNA
 NM_0010016: Homo sapiens FLJ41603 protein (FLJ41603), mRNA
 NM_0010016: Homo sapiens FLJ46321 protein (FLJ46321), mRNA
 NM_0010016: Homo sapiens FLJ16518 protein (FLJ16518), mRNA
 NM_0010016: Homo sapiens ribonuclease-like protein 9 (h461), mRNA

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NM_0010016: Homo sapiens FLJ46385 protein (FLJ46385), mRNA
NM_0010016: Homo sapiens lipocalin 9 (LCN9), mRNA
NM_0010016: Homo sapiens FLJ46300 protein (FLJ46300), mRNA
NM_0010016: Homo sapiens FLJ44653 protein (FLJ44653), mRNA
NM_0010016: Homo sapiens FLJ41423 protein (FLJ41423), mRNA
NM_0010016: Homo sapiens FLJ42102 protein (FLJ42102), mRNA
NM_0010016: Homo sapiens FLJ45300 protein (FLJ45300), mRNA
NM_0010016: Homo sapiens FLJ45530 protein (FLJ45530), mRNA
NM_0010016: Homo sapiens similar to HSPC296 (MGC88387), mRNA
NM_0010016: Homo sapiens FLJ45831 protein (FLJ45831), mRNA
NM_0010016: Homo sapiens FLJ45079 protein (FLJ45079), mRNA
NM_0010016: Homo sapiens FLJ43870 protein (FLJ43870), mRNA
NM_0010016: Homo sapiens FLJ26850 protein (FLJ26850), mRNA
NM_0010016: Homo sapiens FLJ35409 protein (FLJ35409), mRNA
NM_0010016: Homo sapiens FLJ44005 protein (FLJ44005), mRNA
NM_0010016: Homo sapiens hypothetical FLJ42133 (FLJ42133), mRNA
NM_0010016: Homo sapiens FLJ44790 protein (FLJ44790), mRNA
NM_0010016: Homo sapiens FLJ45139 protein (FLJ45139), mRNA
NM_0010016: Homo sapiens FLJ46257 protein (FLJ46257), mRNA
NM_0010016: Homo sapiens FLJ41993 protein (FLJ41993), mRNA
NM_0010016: Homo sapiens FLJ42418 protein (FLJ42418), mRNA
NM_0010016: Homo sapiens FLJ44006 protein (FLJ44006), mRNA
NM_0010016: Homo sapiens FLJ41821 protein (FLJ41821), mRNA
NM_0010016: Homo sapiens FLJ43879 protein (FLJ43879), mRNA
NM_0010016: Homo sapiens FLJ25996 protein (FLJ25996), mRNA
NM_0010017: Homo sapiens FLJ45966 protein (FLJ45966), mRNA
NM_0010017: Homo sapiens HCV F-transactivated protein 1 (LOC401152), mRNA
NM_0010017: Homo sapiens FLJ33360 protein (FLJ33360), mRNA
NM_0010017: Homo sapiens FLJ46010 protein (FLJ46010), mRNA
NM_0010017: Homo sapiens FLJ44796 protein (FLJ44796), mRNA
NM_0010017: Homo sapiens FLJ41649 protein (FLJ41649), mRNA
NM_0010017: Homo sapiens FLJ42177 protein (FLJ42177), mRNA
NM_0010017: Homo sapiens FLJ45974 protein (FLJ45974), mRNA
NM_0010017: Homo sapiens FLJ45537 protein (FLJ45537), mRNA
NM_0010017: Homo sapiens similar to 4931415M17 protein (LOC401565), mRNA
NM_0010017: Homo sapiens DNA-damage inducible protein 1 (DDI1), mRNA
NM_0010017: Homo sapiens lipocalin 10 (LCN10), mRNA
NM_0010017: Homo sapiens SH3 domain binding glutamic acid-rich protein (SH3BGR), tra
NM_0010017: Homo sapiens FERM, RhoGEF (ARHGEF) and pleckstrin domain protein 1 (t
NM_0010017: Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-c
NM_0010017: Homo sapiens chromodomain protein, Y chromosome, 2 related (CDY), mR
NM_0010017: Homo sapiens transmembrane protein 1 (TMEM1), transcript variant 2, mRN
NM_0010017: Homo sapiens Mahlavu hepatocellular carcinoma (HHCm), mRNA
NM_0010017: Homo sapiens chromosome 10 open reading frame 130 (C10orf130), mRNA
NM_0010017: Homo sapiens ATPase, Na⁺/K⁺ transporting, alpha 4 polypeptide (ATP1A4), mR
NM_0010017: Homo sapiens constitutive photomorphogenic protein (COP1), transcript vari
NM_0010017: Homo sapiens BRCC2 mRNA (BRCC2), mRNA
NM_0010017: Homo sapiens ATPase, Na⁺/K⁺ transporting, beta 1 polypeptide (ATP1B1), t
NM_0010017: Homo sapiens chromosome 21 open reading frame 24 (C21orf24), mRNA
NM_0010017: Homo sapiens chromosome 9 open reading frame 105 (C9orf105), mRNA
NM_0010017: Homo sapiens chromosome 10 open reading frame 55 (C10orf55), mRNA
NM_0010017: Homo sapiens similar to RIKEN cDNA 1700027J05 gene (MGC33892), mRN
NM_0010017: Homo sapiens similar to RIKEN cDNA C030006K11 gene (MGC70857), mR
NM_0010018: Homo sapiens olfactory receptor, family 2, subfamily A, member 42 (OR2A4;
NM_0010018: Homo sapiens olfactory receptor, family 2, subfamily T, member 34 (OR2T34;
NM_0010018: Homo sapiens olfactory receptor, family 2, subfamily T, member 27 (OR2T27;
NM_0010018: Homo sapiens olfactory receptor, family 2, subfamily T, member 35 (OR2T35

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NM_0010018: Homo sapiens similar to RIKEN cDNA A030009B12 gene (MGC21382), mRNA
 NM_0010018: Homo sapiens inter-alpha (globulin) inhibitor H5 (ITIH5), transcript variant 3,
 NM_0010018: Homo sapiens serine/threonine-protein kinase pim-3 (PIM3), mRNA
 NM_0010018: Homo sapiens hypothetical protein MGC14376 (MGC14376), transcript varia
 NM_0010018: Homo sapiens heat shock transcription factor, Y-linked 1 (HSFY1), transcript
 NM_0010018: Homo sapiens chromosome 14 open reading frame 37 (C14orf37), mRNA
 NM_0010018: Homo sapiens hypothetical protein LOC283174 (LOC283174), mRNA
 NM_0010018: Homo sapiens protein kinase NYD-SP25 (NYD-SP25), transcript variant 2, r
 NM_0010018: Homo sapiens protein kinase NYD-SP25 (NYD-SP25), transcript variant 3, r
 NM_0010018: Homo sapiens heat shock transcription factor, Y linked 2 (HSFY2), transcript
 NM_0010018: Homo sapiens heat shock transcription factor, Y linked 2 (HSFY2), transcript
 NM_0010018: Homo sapiens interferon-induced protein with tetratricopeptide repeats 1 (IFT
 NM_0010018: Homo sapiens variably charged X-C (VCX-C), mRNA
 NM_0010018: Homo sapiens runt-related transcription factor 1 (acute myeloid leukemia 1; z
 NM_0010018: Homo sapiens prostate cancer associated protein 5 (PCANAP5), transcript v
 NM_0010018: Homo sapiens tetratricopeptide repeat domain 3 (TTC3), transcript variant 2,
 NM_0010018: Homo sapiens ubiquitin associated and SH3 domain containing, A (UBASH3
 NM_0010019: Homo sapiens olfactory receptor, family 4, subfamily E, member 2 (OR4E2),
 NM_0010019: Homo sapiens olfactory receptor, family 52, subfamily N, member 1 (OR52N'
 NM_0010019: Homo sapiens olfactory receptor, family 2, subfamily G, member 3 (OR2G3),
 NM_0010019: Homo sapiens olfactory receptor, family 2, subfamily G, member 2 (OR2G2),
 NM_0010019: Homo sapiens olfactory receptor, family 52, subfamily J, member 3 (OR52J3
 NM_0010019: Homo sapiens olfactory receptor, family 56, subfamily A, member 1 (OR56A'
 NM_0010019: Homo sapiens olfactory receptor, family 5, subfamily BF, member 1 (OR5BF'
 NM_0010019: Homo sapiens olfactory receptor, family 5, subfamily AS, member 1 (OR5AS
 NM_0010019: Homo sapiens olfactory receptor OR11-62 (OR11-62), mRNA
 NM_0010019: Homo sapiens mitochondrial tumor suppressor 1 (MTUS1), transcript variant
 NM_0010019: Homo sapiens mitochondrial tumor suppressor 1 (MTUS1), transcript variant
 NM_0010019: Homo sapiens mitochondrial tumor suppressor 1 (MTUS1), transcript variant
 NM_0010019: Homo sapiens peroxisome proliferative activated receptor, alpha (PPARA), tr
 NM_0010019: Homo sapiens peroxisome proliferative activated receptor, alpha (PPARA), tr
 NM_0010019: Homo sapiens peroxisome proliferative activated receptor, alpha (PPARA), tr
 NM_0010019: Homo sapiens mitochondrial tumor suppressor 1 (MTUS1), transcript variant
 NM_0010019: Homo sapiens LIM homeobox 8 (LHX8), mRNA
 NM_0010019: Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, al
 NM_0010019: Homo sapiens KIAA1914 (KIAA1914), transcript variant 1, mRNA
 NM_0010019: Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, al
 NM_0010019: Homo sapiens chromosome 9 open reading frame 47 (C9orf47), mRNA
 NM_0010019: Homo sapiens 6-pyruvoyl-tetrahydropterin synthase/dimerization cofactor of l
 NM_0010019: Homo sapiens olfactory receptor, family 10, subfamily G, member 9 (OR10G
 NM_0010019: Homo sapiens olfactory receptor, family 5, subfamily A, member 2 (OR5A2),
 NM_0010019: Homo sapiens olfactory receptor, family 13, subfamily C, member 9 (OR13C
 NM_0010019: Homo sapiens olfactory receptor, family 2, subfamily W, member 3 (OR2W3'
 NM_0010019: Homo sapiens olfactory receptor, family 7, subfamily G, member 3 (OR7G3),
 NM_0010019: Homo sapiens olfactory receptor, family 11, subfamily L, member 1 (OR11L1
 NM_0010019: Homo sapiens olfactory receptor, family 5, subfamily W, member 2 (OR5W2'
 NM_0010019: Homo sapiens olfactory receptor, family 13, subfamily C, member 3 (OR13C
 NM_0010019: Homo sapiens olfactory receptor, family 6, subfamily B, member 2 pseudoge
 NM_0010019: Homo sapiens olfactory receptor, family 2, subfamily L, member 8 (OR2L8), i
 NM_0010019: Homo sapiens olfactory receptor, family 2, subfamily T, member 11 (OR2T11
 NM_0010019: Homo sapiens olfactory receptor, family 4, subfamily D, member 5 (OR4D5),
 NM_0010019: Homo sapiens olfactory receptor, family 5, subfamily AT, member 1 (OR5AT'
 NM_0010019: Homo sapiens olfactory receptor, family 5, subfamily D, member 13 (OR5D1'
 NM_0010019: Homo sapiens olfactory receptor, family 6, subfamily S, member 1 (OR6S1),
 NM_0010019: Homo sapiens family with sequence similarity 13, member C1 (FAM13C1), tr
 NM_0010019: Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, ga
 NM_0010019: Homo sapiens pleckstrin homology domain containing, family A (phosphoino

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NM_0010019 Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, de
NM_0010019 Homo sapiens arginyltransferase 1 (ATE1), transcript variant 1, mRNA
NM_0010019 Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, ep
NM_0010019 Homo sapiens regeneration associated muscle protease (DKFZP586H2123)
NM_0010019 Homo sapiens ubiquitin specific protease 16 (USP16), transcript variant 2, m
NM_0010019 Homo sapiens glycoprotein M6B (GPM6B), transcript variant 4, mRNA
NM_0010019 Homo sapiens glycoprotein M6B (GPM6B), transcript variant 1, mRNA
NM_0010019 Homo sapiens glycoprotein M6B (GPM6B), transcript variant 2, mRNA
NM_0010019 Homo sapiens exosome component 10 (EXOSC10), transcript variant 1, mR
NM_0010020 Homo sapiens guanosine monophosphate reductase 2 (GMPR2), transcript
NM_0010020 Homo sapiens guanosine monophosphate reductase 2 (GMPR2), transcript
NM_0010020 Homo sapiens guanosine monophosphate reductase 2 (GMPR2), transcript
NM_0010020 Homo sapiens 5'-nucleotidase, cytosolic IB (NT5C1B), transcript variant 1, m
NM_0010020 Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex, su
NM_0010020 Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex, su
NM_0010020 Homo sapiens host cell factor C1 regulator 1 (XPO1 dependant) (HCFC1R1)
NM_0010020 Homo sapiens host cell factor C1 regulator 1 (XPO1 dependant) (HCFC1R1)
NM_0010020 Homo sapiens phosphofructokinase, liver (PFKL), transcript variant 1, mRNA
NM_0010020 Homo sapiens claudin 18 (CLDN18), transcript variant 2, mRNA
NM_0010020 Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex, su
NM_0010020 Homo sapiens complement component 4B, centromeric (C4B), mRNA
NM_0010020 Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex, su
NM_0010020 Homo sapiens hematological and neurological expressed 1 (HN1), transcript
NM_0010020 Homo sapiens hematological and neurological expressed 1 (HN1), transcript
NM_0010020 Homo sapiens defensin, beta 108 (DEFB108), mRNA
NM_0010020 Homo sapiens astacin-like metalloendopeptidase (M12 family) (ASTL), mRN
NM_0010022 Homo sapiens kallikrein 2, prostatic (KLK2), transcript variant 2, mRNA
NM_0010022 Homo sapiens kallikrein 2, prostatic (KLK2), transcript variant 3, mRNA
NM_0010022 Homo sapiens RAB11 family interacting protein 1 (class I) (RAB11FIP1), trar
NM_0010022 Homo sapiens sodium channel modifier 1 (SCNM1), transcript variant 2, mR
NM_0010022 Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1 anti
NM_0010022 Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1 anti
NM_0010022 Homo sapiens afilipilin protein (AFIPIHILIN), transcript variant 3, mRNA
NM_0010022 Homo sapiens APC11 anaphase promoting complex subunit 11 homolog (ye
NM_0010022 Homo sapiens APC11 anaphase promoting complex subunit 11 homolog (ye
NM_0010022 Homo sapiens APC11 anaphase promoting complex subunit 11 homolog (ye
NM_0010022 Homo sapiens APC11 anaphase promoting complex subunit 11 homolog (ye
NM_0010022 Homo sapiens APC11 anaphase promoting complex subunit 11 homolog (ye
NM_0010022 Homo sapiens ADP-ribosylation-like factor 6 interacting protein 4 (ARL6IP4),
NM_0010022 Homo sapiens ADP-ribosylation-like factor 6 interacting protein 4 (ARL6IP4),
NM_0010022 Homo sapiens SMT3 suppressor of mif two 3 homolog 4 (yeast) (SUMO4), r
NM_0010022 Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex, su
NM_0010022 Homo sapiens acyl-CoA:lysocardiolipin acyltransferase 1 (ALCAT1), transcrip
NM_0010022 Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex, su
NM_0010022 Homo sapiens C1q domain containing 1 (C1QDC1), transcript variant 1, mR
NM_0010022 Homo sapiens chromosome 9 open reading frame 58 (C9orf58), transcript ve
NM_0010022 Homo sapiens zinc finger, FYVE domain containing 27 (ZFVE27), transcrip
NM_0010022 Homo sapiens zinc finger, FYVE domain containing 27 (ZFVE27), transcrip
NM_0010022 Homo sapiens epithelial stromal interaction 1 (breast) (EPSTI1), mRNA
NM_0010022 Homo sapiens c-mir, cellular modulator of immune recognition (MIR), transer
NM_0010022 Homo sapiens c-mir, cellular modulator of immune recognition (MIR), transer
NM_0010022 Homo sapiens exosome component 3 (EXOSC3), transcript variant 2, mRNA
NM_0010022 Homo sapiens Fc fragment of IgG, low affinity IIb, receptor for (CD32) (FCGF
NM_0010022 Homo sapiens Fc fragment of IgG, low affinity IIb, receptor for (CD32) (FCGF
NM_0010022 Homo sapiens Fc fragment of IgG, low affinity IIb, receptor for (CD32) (FCGF
NM_0010022 Homo sapiens putative NFkB activating protein 373 (FLJ23091), transcript ve

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NM_0010022: Homo sapiens flavin containing monooxygenase 3 (FMO3), transcript variant NM_0010022: Homo sapiens GATA binding protein 3 (GATA3), transcript variant 1, mRNA
NM_0010022: Homo sapiens golgi autoantigen, golgin subfamily a, 7 (GOLGA7), transcript NM_0010027: Homo sapiens HIRA interacting protein 5 (HIRIP5), transcript variant 2, mRNA
NM_0010027: Homo sapiens HIRA interacting protein 5 (HIRIP5), transcript variant 3, mRNA
NM_0010027: Homo sapiens HIRA interacting protein 5 (HIRIP5), transcript variant 4, mRNA
NM_0010027: Homo sapiens PTPN13-like, Y-linked, centromeric (PRY), mRNA
NM_0010027: Homo sapiens chromosome 10 open reading frame 78 (C10orf78), transcript NM_0010027: Homo sapiens basic charge, Y-linked, 2 (BPY2), mRNA
NM_0010027: Homo sapiens basic charge, Y-linked, 2 (BPY2), mRNA
NM_0010027: Homo sapiens DnaJ (Hsp40) homolog, subfamily B, member 12 (DNAJB12), NM_0010027: Homo sapiens multiple C2-domains with two transmembrane regions 1 (MC1 NM_0010027: Homo sapiens SMC4 structural maintenance of chromosomes 4-like 1 (yeast NM_0010028: Homo sapiens SMC4 structural maintenance of chromosomes 4-like 1 (yeast NM_0010028: Homo sapiens phosphodiesterase 4D interacting protein (myomegalin) (PDE NM_0010028: Homo sapiens phosphodiesterase 4D interacting protein (myomegalin) (PDE NM_0010028: Homo sapiens phosphodiesterase 4D interacting protein (myomegalin) (PDE NM_0010028: Homo sapiens RAB11 family interacting protein 1 (class I) (RAB11FIP1), trar NM_0010028: Homo sapiens hypothetical protein LOC126208 (LOC126208), mRNA
NM_0010028: Homo sapiens phosphatidylinositol (4,5) bisphosphate 5-phosphatase, A (PII NM_0010028: Homo sapiens protein kinase, lysine deficient 3 (PRKWNK3), transcript varia NM_0010028: Homo sapiens DKFZp434A0131 protein (DKFZP434A0131), transcript varian NM_0010028: Homo sapiens myosin, light polypeptide 4, alkali; atrial, embryonic (MYL4), tr NM_0010028: Homo sapiens suppressor of hairy wing homolog 4 (Drosophila) (SUHW4), tr NM_0010028: Homo sapiens suppressor of hairy wing homolog 4 (Drosophila) (SUHW4), tr NM_0010028: Homo sapiens suppressor of hairy wing homolog 4 (Drosophila) (SUHW4), tr NM_0010028: Homo sapiens CTF8, chromosome transmission fidelity factor 8 homolog (S. NM_0010028: Homo sapiens lymphocyte antigen 6 complex, locus G5C (LY6G5C), transcr NM_0010028: Homo sapiens lymphocyte antigen 6 complex, locus G5C (LY6G5C), transcr NM_0010028: Homo sapiens annexin A2 (ANXA2), transcript variant 2, mRNA
NM_0010028: Homo sapiens annexin A2 (ANXA2), transcript variant 1, mRNA
NM_0010028: Homo sapiens BTB (POZ) domain containing 7 (BTBD7), transcript variant 1. NM_0010028: Homo sapiens Rho guanine nucleotide exchange factor (GEF) 5 (ARHGGEF5 NM_0010028: Homo sapiens chromosome 22 open reading frame 14 (C22orf14), transcript NM_0010028: Homo sapiens chromosome 22 open reading frame 18 (C22orf18), transcript NM_0010028: Homo sapiens chromosome 22 open reading frame 19 (C22orf19), transcript NM_0010028: Homo sapiens chromosome 22 open reading frame 19 (C22orf19), transcript NM_0010028: Homo sapiens chromosome 22 open reading frame 19 (C22orf19), transcript NM_0010028: Homo sapiens chromosome 22 open reading frame 2 (C22orf2), transcript vs NM_0010028: Homo sapiens phosphatidylinositol-3-phosphate/phosphatidylinositol 5-kinas NM_0010029: Homo sapiens hypothetical protein FLJ31052 (FLJ31052), mRNA
NM_0010029: Homo sapiens olfactory receptor, family 8, subfamily G, member 1 (OR8G1P NM_0010029: Homo sapiens X Kell blood group precursor-related, Y-linked 2 (XKRY2), mR NM_0010029: Homo sapiens olfactory receptor, family 8, subfamily K, member 1 (OR8K1), NM_0010029: Homo sapiens KIAA0553 protein (KIAA0553), mRNA
NM_0010029: Homo sapiens cytochrome P450, family 2, subfamily D, polypeptide 9 pseud NM_0010029: Homo sapiens G protein-coupled receptor 139 (GPR139), mRNA
NM_0010029: Homo sapiens chromosome 9 open reading frame 115 (C9orf115), mRNA
NM_0010029: Homo sapiens potassium channel tetramerisation domain containing 11 (KC NM_0010029: Homo sapiens insulin growth factor-like family member 2 (IGFL2), mRNA
NM_0010029: Homo sapiens H2B histone family, member W, testis-specific (H2BFWT), mF NM_0010029: Homo sapiens olfactory receptor, family 8, subfamily D, member 1 (OR8D1), NM_0010029: Homo sapiens olfactory receptor, family 8, subfamily D, member 2 (OR8D2), NM_0010029: Homo sapiens hypothetical protein LOC285016 (LOC285016), mRNA
NM_0010029: Homo sapiens protein expressed in prostate, ovary, testis, and placenta 8 (P NM_0010029: Homo sapiens adenylate kinase 3-like 2 (AK3L2), mRNA
NM_0010029: Homo sapiens similar to PM5 (FLJ43542), mRNA

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NM_0010029: Homo sapiens insulin growth factor-like family member 4 (IGFL4), mRNA
 NM_0010029: Homo sapiens adaptor-related protein complex 3, sigma 1 subunit (AP3S1),
 NM_0010029: Homo sapiens olfactory receptor, family 5, subfamily AP, member 2 (OR5AP
 NM_0010029: Homo sapiens TWIST neighbor (TWISTNB), mRNA
 NM_0010033: Homo sapiens Bicaudal D homolog 1 (Drosophila) (BICD1), transcript variant
 NM_0010033: Homo sapiens DKFZp451A211 protein (DKFZp451A211), mRNA
 NM_0010034: Homo sapiens calcium channel, voltage-dependent, alpha 1I subunit (CACN.
 NM_0010034: Homo sapiens olfactory receptor, family 56, subfamily A, member 3 (OR56A.
 NM_0010036: Homo sapiens protein phosphatase 2, regulatory subunit B, delta isoform (P.
 NM_0010036: Homo sapiens similar to hypothetical protein Y97E10AL1 (DKFZp761P211),
 NM_0010036: Homo sapiens chromosome 18 open reading frame 1 (C18orf1), transcript ve
 NM_0010036: Homo sapiens chromosome 18 open reading frame 1 (C18orf1), transcript ve
 NM_0010036: Homo sapiens leptin receptor (LEPR), transcript variant 2, mRNA
 NM_0010036: Homo sapiens leptin receptor (LEPR), transcript variant 3, mRNA
 NM_0010079: Homo sapiens zeta-chain (TCR) associated protein kinase 70kDa (ZAP70), t
 NM_0011132: Homo sapiens AFG3 ATPase family gene 3-like 1 (yeast) (AFG3L1), mRNA
 NM_0011222: Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
 NM_001369: Homo sapiens dynein, axonemal, heavy polypeptide 5 (DNAH5), mRNA
 NM_001376: Homo sapiens dynein, cytoplasmic, heavy polypeptide 1 (DNCH1), mRNA
 NM_001378: Homo sapiens dynein, cytoplasmic, intermediate polypeptide 2 (DNC12), mR
 NM_001410: Homo sapiens EGF-like-domain, multiple 4 (EGFL4), mRNA
 NM_001547: Homo sapiens interferon-induced protein with tetratricopeptide repeats 2 (IFI
 NM_001556: Homo sapiens inhibitor of kappa light polypeptide gene enhancer in B-cells, I
 NM_001636: Homo sapiens solute carrier family 25 (mitochondrial carrier; adenine nucleo
 NM_001763: Homo sapiens CD1A antigen, a polypeptide (CD1A), mRNA
 NM_001810: Homo sapiens centromere protein B, 80kDa (CENPB), mRNA
 NM_001931: Homo sapiens dihydrolipoamide S-acetyltransferase (E2 component of pyruv
 NM_001947: Homo sapiens dual specificity phosphatase 7 (DUSP7), mRNA
 NM_001984: Homo sapiens esterase D/formylglutathione hydrolase (ESD), mRNA
 NM_001986: Homo sapiens ets variant gene 4 (E1A enhancer binding protein, E1AF) (ET
 NM_002154: Homo sapiens heat shock 70kDa protein 4 (HSPA4), transcript variant 1, mR
 NM_002242: Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 1
 NM_002348: Homo sapiens lymphocyte antigen 9 (LY9), mRNA
 NM_002399: Homo sapiens Meis1, myeloid ecotropic viral integration site 1 homolog 2 (m
 NM_002404: Homo sapiens microfibrillar-associated protein 4 (MFAP4), mRNA
 NM_002471: Homo sapiens myosin, heavy polypeptide 6, cardiac muscle, alpha (cardiom
 NM_002498: Homo sapiens NIMA (never in mitosis gene a)-related kinase 3 (NEK3), tran
 NM_002523: Homo sapiens neuronal pentraxin II (NPTX2), mRNA
 NM_002596: Homo sapiens PCTAIRE protein kinase 3 (PCTK3), transcript variant 3, mR
 NM_002803: Homo sapiens phosphodiesterase 7A (PDE7A), transcript variant 1, mRNA
 NM_002604: Homo sapiens phosphodiesterase 7A (PDE7A), transcript variant 2, mRNA
 NM_002605: Homo sapiens phosphodiesterase 8A (PDE8A), transcript variant 1, mRNA
 NM_002679: Homo sapiens postmeiotic segregation increased 2-like 2 (PMS2L2), mRNA
 NM_002735: Homo sapiens protein kinase, cAMP-dependent, regulatory, type I, beta (PRI
 NM_002746: Homo sapiens mitogen-activated protein kinase 3 (MAPK3), mRNA
 NM_002791: Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 6 (PSI
 NM_002798: Homo sapiens proteasome (prosome, macropain) subunit, beta type, 6 (PSI
 NM_002972: Homo sapiens SET binding factor 1 (SBF1), transcript variant 1, mRNA
 NM_002974: Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin),
 NM_002988: Homo sapiens syndecan 2 (heparan sulfate proteoglycan 1, cell surface-ass
 NM_003013: Homo sapiens secreted frizzled-related protein 2 (SFRP2), mRNA
 NM_003047: Homo sapiens solute carrier family 9 (sodium/hydrogen exchanger), isoform
 NM_003106: Homo sapiens SRY (sex determining region Y)-box 2 (SOX2), mRNA
 NM_003111: Homo sapiens Sp3 transcription factor (SP3), mRNA
 NM_003179: Homo sapiens synapophysin (SYP), mRNA
 NM_003196: Homo sapiens transcription elongation factor A (SII), 3 (TCEA3), mRNA
 NM_003200: Homo sapiens transcription factor 3 (E2A immunoglobulin enhancer binding

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NM_003302 Homo sapiens thyroid hormone receptor interactor 6 (TRIP6), mRNA
 NM_003415 Homo sapiens zinc finger protein 268 (ZNF268), mRNA
 NM_003444 Homo sapiens zinc finger protein 154 (pH2-92) (ZNF154), mRNA
 NM_003502 Homo sapiens axin 1 (AXIN1), transcript variant 1, mRNA
 NM_003517 Homo sapiens histone 2, H2ac (HIST2H2AC), mRNA
 NM_003575 Homo sapiens zinc finger protein 282 (ZNF282), mRNA
 NM_003598 Homo sapiens TEA domain family member 2 (TEAD2), mRNA
 NM_003638 Homo sapiens integrin, alpha 8 (ITGA8), mRNA
 NM_003660 Homo sapiens protein tyrosine phosphatase, receptor type, f polypeptide (PT
 NM_003677 Homo sapiens density-regulated protein (DENR), mRNA
 NM_003700 Homo sapiens olfactory receptor, family 2, subfamily D, member 2 (OR2D2),
 NM_003719 Homo sapiens phosphodiesterase 8B (PDE8B), mRNA
 NM_003724 Homo sapiens jerky homolog (mouse) (JRK), mRNA
 NM_003741 Homo sapiens chordin (CHRD), transcript variant 1, mRNA
 NM_003817 Homo sapiens a disintegrin and metalloproteinase domain 7 (ADAM7), mRNA
 NM_003818 Homo sapiens CDP-diacylglycerol synthase (phosphatidate cytidyltransferase
 NM_003828 Homo sapiens myotubularin related protein 1 (MTMR1), transcript variant 1,
 NM_003845 Homo sapiens dual-specificity tyrosine-(Y)-phosphorylation regulated kinase
 NM_003848 Homo sapiens succinate-CoA ligase, GDP-forming, beta subunit (SUCLG2),
 NM_003858 Homo sapiens cyclin K (CCNK), mRNA
 NM_003898 Homo sapiens synaptotagmin 2 (SYNJ2), mRNA
 NM_003907 Homo sapiens eukaryotic translation initiation factor 2B, subunit 5 epsilon, 8'
 NM_003957 Homo sapiens serine/threonine kinase 29 (STK29), mRNA
 NM_003959 Homo sapiens huntingtin interacting protein-1-related (HIP1R), mRNA
 NM_003972 Homo sapiens BTAf1 RNA polymerase II, B-TFIID transcription factor-associ
 NM_004080 Homo sapiens diacylglycerol kinase, beta 90kDa (DGKB), transcript variant 1
 NM_004097 Homo sapiens empty spiracles homolog 1 (Drosophila) (EMX1), mRNA
 NM_004118 Homo sapiens forkhead-like 18 (Drosophila) (FKHL18), mRNA
 NM_004136 Homo sapiens Iron-responsive element binding protein 2 (IREB2), mRNA
 NM_004200 Homo sapiens synaptotagmin VII (SYT7), mRNA
 NM_004220 Homo sapiens zinc finger protein 213 (ZNF213), mRNA
 NM_004241 Homo sapiens jumonji domain containing 1C (JMJD1C), mRNA
 NM_004242 Homo sapiens high mobility group nucleosomal binding domain 3 (HMGN3),
 NM_004319 Homo sapiens astrotactin (ASTN), transcript variant 1, mRNA
 NM_004439 Homo sapiens EphA5 (EPHA5), transcript variant 1, mRNA
 NM_004498 Homo sapiens one cut domain, family member 1 (ONECUT1), mRNA
 NM_004650 Homo sapiens GS2 gene (DXS1283E), mRNA
 NM_004686 Homo sapiens myotubularin related protein 6 (MTMR6), mRNA
 NM_004691 Homo sapiens ATPase, H⁺ transporting, lysosomal 38kDa, V0 subunit d isofo
 NM_004764 Homo sapiens piwi-like 1 (Drosophila) (PIWIL1), mRNA
 NM_004773 Homo sapiens thyroid hormone receptor interactor 3 (TRIP3), mRNA
 NM_004816 Homo sapiens chromosome 9 open reading frame 61 (C9orf61), mRNA
 NM_004840 Homo sapiens Rac/Cdc42 guanine nucleotide exchange factor (GEF) 6 (ARF
 NM_004884 Homo sapiens putative neuronal cell adhesion molecule (PUNC), mRNA
 NM_004946 Homo sapiens dedicator of cytokinesis 2 (DOCK2), mRNA
 NM_004947 Homo sapiens dedicator of cytokinesis 3 (DOCK3), mRNA
 NM_005054 Homo sapiens RAN binding protein 2-like 1 (RANBP2L1), transcript variant 1
 NM_005105 Homo sapiens RNA binding motif protein 8A (RBM8A), mRNA
 NM_005126 Homo sapiens nuclear receptor subfamily 1, group D, member 2 (NR1D2), tr
 NM_005140 Homo sapiens cyclic nucleotide gated channel alpha 2 (CNGA2), mRNA
 NM_005144 Homo sapiens hairless homolog (mouse) (HR), transcript variant 1, mRNA
 NM_005153 Homo sapiens ubiquitin specific protease 10 (USP10), mRNA
 NM_005202 Homo sapiens collagen, type VIII, alpha 2 (COL8A2), mRNA
 NM_005240 Homo sapiens ets variant gene 3 (ETV3), mRNA
 NM_005241 Homo sapiens ecotropic viral integration site 1 (EVI1), mRNA
 NM_005250 Homo sapiens forkhead box L1 (FOXL1), mRNA
 NM_005272 Homo sapiens guanine nucleotide binding protein (G protein), alpha transduc

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NM_005278 Homo sapiens glycoprotein M6B (GPM6B), transcript variant 3, mRNA
 NM_005349 Homo sapiens recombining binding protein suppressor of hairless (Drosophila)
 NM_005376 Homo sapiens v-myc myelocytomatosis viral oncogene homolog 1, lung carc
 NM_005407 Homo sapiens sal-like 2 (Drosophila) (SALL2), mRNA
 NM_005482 Homo sapiens phosphatidylinositol glycan, class K (PIGK), mRNA
 NM_005487 Homo sapiens high-mobility group protein 2-like 1 (HMG2L1), mRNA
 NM_005533 Homo sapiens interferon-induced protein 35 (IFI35), mRNA
 NM_005559 Homo sapiens laminin, alpha 1 (LAMA1), mRNA
 NM_005595 Homo sapiens nuclear factor I/A (NFIA), mRNA
 NM_005650 Homo sapiens transcription factor 20 (AR1) (TCF20), transcript variant 1, mF
 NM_005669 Homo sapiens chromosome 5 open reading frame 18 (C5orf18), mRNA
 NM_005680 Homo sapiens TATA box binding protein (TBP)-associated factor, RNA polyn
 NM_005702 Homo sapiens Era G-protein-like 1 (E. coli) (ERAL1), mRNA
 NM_005707 Homo sapiens programmed cell death 7 (PDCD7), mRNA
 NM_005779 Homo sapiens lipoma HMGIC fusion partner-like 2 (LHFPL2), mRNA
 NM_005788 Homo sapiens HMT1 hnRNP methyltransferase-like 3 (S. cerevisiae) (HRMT
 NM_005791 Homo sapiens M-phase phosphoprotein 10 (U3 small nucleolar ribonucleopr
 NM_005840 Homo sapiens sprouty homolog 3 (Drosophila) (SPRY3), mRNA
 NM_005841 Homo sapiens sprouty homolog 1, antagonist of FGF signaling (Drosophila) i
 NM_005848 Homo sapiens c-myc promoter binding protein (MYCPBP), mRNA
 NM_005914 Homo sapiens MCM4 minichromosome maintenance deficient 4 (S. cerevisie
 NM_005942 Homo sapiens molybdenum cofactor synthesis 1 (MOCOS1), transcript variant
 NM_005943 Homo sapiens molybdenum cofactor synthesis 1 (MOCOS1), transcript variant
 NM_005946 Homo sapiens metallothionein 1A (functional) (MT1A), mRNA
 NM_005947 Homo sapiens metallothionein 1B (functional) (MT1B), mRNA
 NM_005949 Homo sapiens metallothionein 1F (functional) (MT1F), mRNA
 NM_005964 Homo sapiens myosin, heavy polypeptide 10, non-muscle (MYH10), mRNA
 NM_005984 Homo sapiens solute carrier family 25 (mitochondrial carrier; citrate transport
 NM_005995 Homo sapiens T-box 10 (TBX10), mRNA
 NM_006036 Homo sapiens putative prolyl oligopeptidase (KIAA0436), mRNA
 NM_006040 Homo sapiens heparan sulfate (glucosamine) 3-O-sulfotransferase 4 (HS3S1
 NM_006082 Homo sapiens SMYD family member 5 (SMYD5), mRNA
 NM_006091 Homo sapiens coronin, actin binding protein, 2B (CORO2B), mRNA
 NM_006108 Homo sapiens spondin 1, extracellular matrix protein (SPON1), mRNA
 NM_006133 Homo sapiens chromosome 11 open reading frame 11 (C11orf11), mRNA
 NM_006151 Homo sapiens lactoperoxidase (LPO), mRNA
 NM_006154 Homo sapiens neural precursor cell expressed, developmentally down-regule
 NM_006172 Homo sapiens natriuretic peptide precursor A (NPPA), mRNA
 NM_006175 Homo sapiens nebulin-related anchoring protein (NRAP), transcript variant 1
 NM_006210 Homo sapiens paternally expressed 3 (PEG3), mRNA
 NM_006216 Homo sapiens serine (or cysteine) proteinase inhibitor, clade E (nexin, plasr
 NM_006266 Homo sapiens rat guanine nucleotide dissociation stimulator (RALGDS), mR
 NM_006277 Homo sapiens intersectin 2 (ITSN2), transcript variant 1, mRNA
 NM_006452 Homo sapiens phosphoribosylaminoimidazole carboxylase, phosphoribosyla
 NM_006524 Homo sapiens zinc finger protein 138 (clone pHZ-32) (ZNF138), mRNA
 NM_006591 Homo sapiens polymerase (DNA-directed), delta 3, accessory subunit (POLC
 NM_006617 Homo sapiens nestin (NES), mRNA
 NM_006630 Homo sapiens zinc finger protein 234 (ZNF234), mRNA
 NM_006631 Homo sapiens zinc finger protein 266 (ZNF266), mRNA
 NM_006635 Homo sapiens zinc finger protein 272 (ZNF272), mRNA
 NM_006642 Homo sapiens serologically defined colon cancer antigen 8 (SDCCAG8), mR
 NM_006647 Homo sapiens NADPH oxidase activator 1 (NOXA1), mRNA
 NM_006673 Homo sapiens AT rich interactive domain 5A (MRF1-like) (ARID5A), transcr
 NM_006714 Homo sapiens sphingomyelin phosphodiesterase, acid-like 3A (SMPDL3A), i
 NM_006722 Homo sapiens microphthalmia-associated transcription factor (MITF), transcr
 NM_006742 Homo sapiens protein serine kinase H1 (PSKH1), mRNA
 NM_006775 Homo sapiens quaking homolog, KH domain RNA binding (mouse) (QKI), trs

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NM_006828 Homo sapiens activating signal cointegrator 1 complex subunit 3 (ASCC3), n
 NM_006832 Homo sapiens pleckstrin homology domain containing, family C (with FERM
 NM_006857 Homo sapiens putative nucleic acid binding protein RY-1 (RY1), mRNA
 NM_006859 Homo sapiens lipic acid synthetase (LIAS), nuclear gene encoding mitoch
 NM_006897 Homo sapiens homeo box C9 (HOXC9), mRNA
 NM_006909 Homo sapiens Ras protein-specific guanine nucleotide-releasing factor 2 (R/
 NM_006916 Homo sapiens ribulose-5-phosphate-3-epimerase (RPE), transcript variant 2.
 NM_006920 Homo sapiens sodium channel, voltage-gated, type I, alpha (SCN1A), mRNA
 NM_006939 Homo sapiens son of sevenless homolog 2 (Drosophila) (SOS2), mRNA
 NM_006955 Homo sapiens zinc finger protein 11b (KOX 2) (ZNF11b), mRNA
 NM_006956 Homo sapiens zinc finger protein 12 (KOX 3) (ZNF12), mRNA
 NM_006959 Homo sapiens zinc finger protein 17 (HPF3, KOX 10) (ZNF17), mRNA
 NM_006961 Homo sapiens zinc finger protein 19 (KOX 12) (ZNF19), mRNA
 NM_006969 Homo sapiens zinc finger protein 28 (KOX 24) (ZNF28), mRNA
 NM_006973 Homo sapiens zinc finger protein 32 (KOX 30) (ZNF32), mRNA
 NM_006974 Homo sapiens zinc finger protein 33a (KOX 31) (ZNF33A), mRNA
 NM_006996 Homo sapiens solute carrier family 19 (thiamine transporter), member 2 (SLC
 NM_007001 Homo sapiens solute carrier family 35, member D2 (SLC35D2), mRNA
 NM_007010 Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 52 (DDX52), transc
 NM_007041 Homo sapiens arginyltransferase 1 (ATE1), transcript variant 2, mRNA
 NM_007078 Homo sapiens LIM domain binding 3 (LDB3), mRNA
 NM_007130 Homo sapiens zinc finger protein 41 (ZNF41), transcript variant 1, mRNA
 NM_007131 Homo sapiens zinc finger protein 75 (D8C6) (ZNF75), mRNA
 NM_007135 Homo sapiens zinc finger protein 79 (p17) (ZNF79), mRNA
 NM_007137 Homo sapiens zinc finger protein 81 (HFZ20) (ZNF81), mRNA
 NM_007139 Homo sapiens zinc finger protein 92 (HTF12) (ZNF92), mRNA
 NM_007149 Homo sapiens zinc finger protein 184 (Kruppel-like) (ZNF184), mRNA
 NM_007156 Homo sapiens zinc finger, X-linked, duplicated A (ZXDA), mRNA
 NM_007157 Homo sapiens zinc finger, X-linked, duplicated B (ZXDB), mRNA
 NM_007162 Homo sapiens transcription factor EB (TFEB), mRNA
 NM_007174 Homo sapiens citron (rho-interacting, serine/threonine kinase 21) (CIT), mR
 NM_007189 Homo sapiens ATP-binding cassette, sub-family F (Gcn20), member 2 (ABC
 NM_007224 Homo sapiens neurexophilin 4 (NXP4), mRNA
 NM_007225 Homo sapiens neurexophilin 3 (NXP3), mRNA
 NM_007243 Homo sapiens nufm (nuclear envelope membrane protein) (NRM), mRNA
 NM_007261 Homo sapiens leukocyte membrane antigen (CMRF-35H), mRNA
 NM_007270 Homo sapiens FK506 binding protein 9, 63 kDa (FKBP9), mRNA
 NM_007277 Homo sapiens SEC6-like 1 (S. cerevisiae) (SEC6L1), mRNA
 NM_007280 Homo sapiens Opa-interacting protein 5 (OIP5), mRNA
 NM_007349 Homo sapiens PAX transcription activation domain interacting protein 1 like (/
 NM_007356 Homo sapiens laminin, beta 4 (LAMB4), mRNA
 NM_012073 Homo sapiens chaperonin containing TCP1, subunit 5 (epsilon) (CCT5), mR
 NM_012154 Homo sapiens eukaryotic translation initiation factor 2C, 2 (EIF2C2), mRNA
 NM_012156 Homo sapiens erythrocyte membrane protein band 4.1-like 1 (EPB41L1), tra
 NM_012167 Homo sapiens F-box protein 11 (FBXO11), transcript variant 3, mRNA
 NM_012174 Homo sapiens F-box and WD-40 domain protein 8 (FBXW8), transcript varia
 NM_012184 Homo sapiens forkhead box D4 like 1 (FOXO4L1), mRNA
 NM_012212 Homo sapiens leukotriene B4 12-hydroxydehydrogenase (LTB4DH), mRNA
 NM_012224 Homo sapiens NIMA (never in mitosis gene a)-related kinase 1 (NEK1), mR
 NM_012232 Homo sapiens polymerase I and transcript release factor (PTRF), mRNA
 NM_012235 Homo sapiens SREBP CLEAVAGE-ACTIVATING PROTEIN (SCAP), mRNA
 NM_012271 Homo sapiens huntingtin interacting protein B (HYPB), transcript variant 2, r
 NM_012272 Homo sapiens Huntingtin interacting protein C (HYPC), mRNA
 NM_012284 Homo sapiens potassium voltage-gated channel, subfamily H (eag-related), i
 NM_012292 Homo sapiens minor histocompatibility antigen HA-1 (HA-1), mRNA
 NM_012305 Homo sapiens adaptor-related protein complex 2, alpha 2 subunit (AP2A2), r
 NM_012309 Homo sapiens SH3 and multiple ankyrin repeat domains 2 (SHANK2), transc

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NM_012315 Homo sapiens kallikrein 9 (KLK9), mRNA
 NM_012335 Homo sapiens myosin IF (MYO1F), mRNA
 NM_012363 Homo sapiens olfactory receptor, family 1, subfamily N, member 1 (OR1N1),
 NM_012364 Homo sapiens olfactory receptor, family 1, subfamily Q, member 1 (OR1Q1),
 NM_012367 Homo sapiens olfactory receptor, family 2, subfamily B, member 6 (OR2B6),
 NM_012374 Homo sapiens olfactory receptor, family 4, subfamily D, member 1 (OR4D1),
 NM_012378 Homo sapiens olfactory receptor, family 8, subfamily B, member 8 (OR8B8),
 NM_012393 Homo sapiens phosphoribosylformylglycinamide synthase (FGAR amidotransferase),
 NM_012398 Homo sapiens phosphatidylinositol 4-phosphate 5-kinase, type I, gamma (PI3Kγ),
 NM_012416 Homo sapiens RAN binding protein 6 (RANBP6), mRNA
 NM_012477 Homo sapiens WW domain binding protein 1 (WBP1), mRNA
 NM_012478 Homo sapiens WW domain binding protein 2 (WBP2), mRNA
 NM_013304 Homo sapiens zinc finger, DHHC domain containing 1 (ZDHHC1), mRNA
 NM_013321 Homo sapiens sorting nexin 8 (SNX8), mRNA
 NM_013373 Homo sapiens zinc finger, DHHC domain containing 8 (ZDHHC8), mRNA
 NM_014010 Homo sapiens astrotactin 2 (ASTN2), transcript variant 1, mRNA
 NM_014014 Homo sapiens U5 snRNP-specific protein, 200-KD (U5-200KD), mRNA
 NM_014089 Homo sapiens nucleoporin like 1 (NUPL1), mRNA
 NM_014098 Homo sapiens peroxiredoxin 3 (PRDX3), nuclear gene encoding mitochondria,
 NM_014215 Homo sapiens insulin receptor-related receptor (INSRR), mRNA
 NM_014220 Homo sapiens transmembrane 4 superfamily member 1 (TM4SF1), mRNA
 NM_014224 Homo sapiens pepsinogen 5, group I (pepsinogen A) (PGA5), mRNA
 NM_014243 Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with thiol
 NM_014261 Homo sapiens TIR domain containing adaptor inducing interferon-beta (TRIF),
 NM_014282 Homo sapiens hyaluronan binding protein 4 (HABP4), mRNA
 NM_014284 Homo sapiens neurochondrin (NCDN), mRNA
 NM_014290 Homo sapiens tudor domain containing 7 (TDRD7), mRNA
 NM_014301 Homo sapiens iron-sulfur cluster assembly enzyme (ISCU), mRNA
 NM_014346 Homo sapiens chromosome 22 open reading frame 4 (C22orf4), mRNA
 NM_014376 Homo sapiens cytoplasmic FMR1 interacting protein 2 (CYFIP2), mRNA
 NM_014381 Homo sapiens mutL homolog 3 (E. coli) (MLH3), mRNA
 NM_014389 Homo sapiens proline-, glutamic acid-, leucine-rich protein 1 (PELP1), mRNA
 NM_014422 Homo sapiens phosphatidylinositol (4,5) bisphosphate 5-phosphatase, A (PLI1),
 NM_014435 Homo sapiens N-acylsphingosine amidohydrolase (acid ceramidase)-like (AC3),
 NM_014441 Homo sapiens sialic acid binding Ig-like lectin 9 (SIGLEC9), mRNA
 NM_014455 Homo sapiens zinc finger protein 364 (ZNF364), mRNA
 NM_014460 Homo sapiens RNA-binding protein p1ppin (PIPPIN), mRNA
 NM_014472 Homo sapiens chromosome 10 open reading frame 28 (C10orf28), mRNA
 NM_014494 Homo sapiens trinucleotide repeat containing 6 (TNRC6), mRNA
 NM_014507 Homo sapiens malonyl-CoA:acyl carrier protein transacylase (malonyltransferase),
 NM_014508 Homo sapiens apolipoprotein B mRNA editing enzyme, catalytic polypeptide,
 NM_014510 Homo sapiens piccolo (presynaptic cytomatrix protein) (PCLC), transcript variant 1,
 NM_014562 Homo sapiens orthodenticle homolog 1 (Drosophila) (OTX1), mRNA
 NM_014568 Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetyltransferase,
 NM_014572 Homo sapiens LATs, large tumor suppressor, homolog 2 (Drosophila) (LATs),
 NM_014573 Homo sapiens hypothetical protein MAC30 (MAC30), mRNA
 NM_014594 Homo sapiens zinc finger protein 354C (ZNF354C), mRNA
 NM_014602 Homo sapiens phosphoinositide-3-kinase, regulatory subunit 4, p150 (PIK3R4),
 NM_014603 Homo sapiens paraneoplastic antigen (HUMPPA), mRNA
 NM_014607 Homo sapiens UBX domain containing 2 (UBXD2), mRNA
 NM_014608 Homo sapiens cytoplasmic FMR1 interacting protein 1 (CYFIP1), mRNA
 NM_014611 Homo sapiens MDN1, midasin homolog (yeast) (MDN1), mRNA
 NM_014613 Homo sapiens expressed in T-cells and eosinophils in atopic dermatitis (ETE),
 NM_014614 Homo sapiens proteasome (prosome, macropain) activator subunit 4 (PSME4),
 NM_014615 Homo sapiens KIAA0182 protein (KIAA0182), mRNA
 NM_014647 Homo sapiens limkain b1 (LKAP), transcript variant 1, mRNA
 NM_014655 Homo sapiens KIAA0446 gene product (KIAA0446), mRNA

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NM_014657 Homo sapiens KIAA0406 gene product (KIAA0406), mRNA
 NM_014667 Homo sapiens vestigial like 4 (Drosophila) (VGLL4), mRNA
 NM_014691 Homo sapiens aquarius homolog (mouse) (AQR), mRNA
 NM_014697 Homo sapiens C-terminal PDZ domain ligand of neuronal nitric oxide synthase
 NM_014701 Homo sapiens KIAA0256 gene product (KIAA0256), mRNA
 NM_014756 Homo sapiens KIAA0097 gene product (ch-TOG), mRNA
 NM_014788 Homo sapiens pleckstrin homology domain containing, family M (with RUN d
 NM_014802 Homo sapiens KIAA0528 gene product (KIAA0528), mRNA
 NM_014836 Homo sapiens Rho-related BTB domain containing 1 (RHOBTB1), transcript
 NM_014839 Homo sapiens plasticity related gene 1 (LPPR4), mRNA
 NM_014850 Homo sapiens SLIT-ROBO Rho GTPase activating protein 2 (SRGAP2), mR
 NM_014854 Homo sapiens solute carrier family 35, member E2 (SLC35E2), mRNA
 NM_014858 Homo sapiens cerebral protein 11 (HUCEP11), mRNA
 NM_014881 Homo sapiens DNA cross-link repair 1A (PSO2 homolog, S. cerevisiae) (DCI
 NM_014884 Homo sapiens splicing factor, arginine/serine-rich 14 (SFRS14), mRNA
 NM_014919 Homo sapiens Wolf-Hirschhorn syndrome candidate 1 (WHSC1), transcript v
 NM_014955 Homo sapiens CGI-01 protein (CGI-01), transcript variant 2, mRNA
 NM_014957 Homo sapiens KIAA0870 protein (KIAA0870), mRNA
 NM_014974 Homo sapiens KIAA0934 (KIAA0934), mRNA
 NM_014975 Homo sapiens microtubule associated serine/threonine kinase 1 (MAST1), m
 NM_014982 Homo sapiens pecanex homolog (Drosophila) (PCNX), mRNA
 NM_014989 Homo sapiens regulating synaptic membrane exocytosis 1 (RIMS1), transcrip
 NM_014991 Homo sapiens WD repeat and FYVE domain containing 3 (WDFY3), transcri
 NM_014992 Homo sapiens dishevelled associated activator of morphogenesis 1 (DAAM1
 NM_014997 Homo sapiens KIAA0265 protein (KIAA0265), mRNA
 NM_015000 Homo sapiens serine/threonine kinase 38 like (STK38L), mRNA
 NM_015004 Homo sapiens exosome component 7 (EXOSC7), mRNA
 NM_015008 Homo sapiens KIAA0779 protein (KIAA0779), mRNA
 NM_015013 Homo sapiens amine oxidase (flavin containing) domain 2 (AOF2), mRNA
 NM_015014 Homo sapiens KIAA0117 protein (KIAA0117), mRNA
 NM_015015 Homo sapiens jumonji domain containing 2B (JMJD2B), mRNA
 NM_015017 Homo sapiens ubiquitin specific protease 33 (USP33), transcript variant 1, m
 NM_015018 Homo sapiens KIAA1117 (KIAA1117), mRNA
 NM_015022 Homo sapiens PDZ domain containing 3 (PDZK3), transcript variant 2, mRN
 NM_015027 Homo sapiens KIAA0251 protein (KIAA0251), mRNA
 NM_015029 Homo sapiens processing of precursor 1, ribonuclease P/MRP subunit (S. ce
 NM_015033 Homo sapiens formin binding protein 1 (FBNP1), mRNA
 NM_015035 Homo sapiens zinc fingers and homeoboxes 3 (ZHX3), mRNA
 NM_015037 Homo sapiens KIAA0913 (KIAA0913), mRNA
 NM_015039 Homo sapiens nicotinamide nucleotide adenyllyltransferase 2 (NMNAT2), tra
 NM_015040 Homo sapiens phosphatidylinositol-3-phosphate/phosphatidylinositol 5-kinas
 NM_015045 Homo sapiens KIAA0261 (KIAA0261), mRNA
 NM_015047 Homo sapiens KIAA0090 protein (KIAA0090), mRNA
 NM_015050 Homo sapiens KIAA0082 (KIAA0082), mRNA
 NM_015052 Homo sapiens HECT type E3 ubiquitin ligase (NEDL1), mRNA
 NM_015055 Homo sapiens SWAP-70 protein (SWAP70), mRNA
 NM_015059 Homo sapiens talin 2 (TLN2), mRNA
 NM_015061 Homo sapiens jumonji domain containing 2C (JMJD2C), mRNA
 NM_015065 Homo sapiens SLAC2-B (SLAC2-B), mRNA
 NM_015066 Homo sapiens tripartite motif-containing 35 (TRIM35), transcript variant 1, m
 NM_015069 Homo sapiens zinc finger protein 423 (ZNF423), mRNA
 NM_015076 Homo sapiens cyclin-dependent kinase (CDC2-like) 11 (CDK11), mRNA
 NM_015078 Homo sapiens Rho family guanine-nucleotide exchange factor (KIAA0861), r
 NM_015079 Homo sapiens KIAA1055 protein (KIAA1055), mRNA
 NM_015085 Homo sapiens GTPase activating RANGAP domain-like 4 (GARNL4), mRNA
 NM_015087 Homo sapiens spastic paraplegia 20, spartin (Troyer syndrome) (SPG20), m
 NM_015089 Homo sapiens p53-associated parkin-like cytoplasmic protein (PARC), mRN

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NM_015091 Homo sapiens KIAA0423 (KIAA0423), mRNA
 NM_015094 Homo sapiens hypermethylated in cancer 2 (HIC2), mRNA
 NM_015099 Homo sapiens calmodulin binding transcription activator 2 (CAMTA2), mRNA
 NM_015100 Homo sapiens pogo transposable element with ZNF domain (POGZ), transcr
 NM_015102 Homo sapiens nephronophthisis 4 (NPHP4), mRNA
 NM_015103 Homo sapiens plexin D1 (PLXND1), mRNA
 NM_015106 Homo sapiens KIAA0809 protein (SRISNF2L), mRNA
 NM_015107 Homo sapiens PHD finger protein 8 (PHF8), mRNA
 NM_015110 Homo sapiens SMC5 structural maintenance of chromosomes 5-like 1 (yeast
 NM_015115 Homo sapiens KIAA0276 protein (KIAA0276), mRNA
 NM_015116 Homo sapiens leucine-rich repeats and calponin homology (CH) domain con
 NM_015117 Homo sapiens zinc finger CCHC type domain containing 3 (ZC3HDC3), mRN
 NM_015120 Homo sapiens Alstrom syndrome 1 (ALMS1), mRNA
 NM_015122 Homo sapiens FCH domain only 1 (FCHO1), mRNA
 NM_015134 Homo sapiens myosin phosphatase-Rho interacting protein (M-RIP), mRNA
 NM_015138 Homo sapiens KIAA0252 (KIAA0252), mRNA
 NM_015141 Homo sapiens glycerol-3-phosphate dehydrogenase 1-like (GPD1L), mRNA
 NM_015143 Homo sapiens methionyl aminopeptidase 1 (METAP1), mRNA
 NM_015144 Homo sapiens zinc finger, CCHC domain containing 14 (ZCCHC14), mRNA
 NM_015150 Homo sapiens raft-linking protein (RAFTLIN), mRNA
 NM_015151 Homo sapiens chromosome 21 open reading frame 106 (C21orf106), transcr
 NM_015157 Homo sapiens pleckstrin homology-like domain, family B, member 1 (PHLDE
 NM_015158 Homo sapiens ankyrin repeat domain 15 (ANKRD15), transcript variant 1, m
 NM_015160 Homo sapiens peptidase (mitochondrial processing) alpha (PMPCA), nuclea
 NM_015161 Homo sapiens ADP-ribosylation factor-like 6 interacting protein (ARL6IP), m
 NM_015167 Homo sapiens phosphatidylserine receptor (PTDSR), mRNA
 NM_015170 Homo sapiens sulfatase 1 (SULF1), mRNA
 NM_015171 Homo sapiens exportin 6 (XPO6), mRNA
 NM_015172 Homo sapiens HBxAg transactivated protein 2 (XTP2), mRNA
 NM_015173 Homo sapiens TBC1 (tre-2/USP6, BUB2, cdc16) domain family, member 1 (T
 NM_015184 Homo sapiens phospholipase C-like 2 (PLCL2), mRNA
 NM_015187 Homo sapiens KIAA0746 protein (KIAA0746), mRNA
 NM_015190 Homo sapiens DnaJ (Hsp40) homolog, subfamily C, member 9 (DNAJC9), m
 NM_015191 Homo sapiens salt-inducible serine/threonine kinase 2 (SIK2), mRNA
 NM_015198 Homo sapiens cordon-bleu homolog (mouse) (COBL), mRNA
 NM_015199 Homo sapiens ankyrin repeat domain 28 (ANKRD28), mRNA
 NM_015200 Homo sapiens KIAA0648 protein (KIAA0648), mRNA
 NM_015201 Homo sapiens block of proliferation 1 (BOP1), mRNA
 NM_015203 Homo sapiens KIAA0460 protein (KIAA0460), mRNA
 NM_015210 Homo sapiens KIAA0802 (KIAA0802), mRNA
 NM_015213 Homo sapiens RAB6 interacting protein 1 (RAB6IP1), mRNA
 NM_015219 Homo sapiens exocyst complex component 7 (EXOC7), mRNA
 NM_015221 Homo sapiens dynamin binding protein (DNMBP), mRNA
 NM_015229 Homo sapiens KIAA0664 protein (KIAA0664), mRNA
 NM_015234 Homo sapiens G protein-coupled receptor 116 (GPR116), mRNA
 NM_015238 Homo sapiens KIBRA protein (KIBRA), mRNA
 NM_015243 Homo sapiens Cohen syndrome 1 (COH1), transcript variant 3, mRNA
 NM_015245 Homo sapiens ankyrin repeat and sterile alpha motif domain containing 1 (A
 NM_015246 Homo sapiens mahogunin, ring finger 1 (MGRN1), mRNA
 NM_015250 Homo sapiens bicaudal D homolog 2 (Drosophila) (BCD2), mRNA
 NM_015252 Homo sapiens NPF/calponin-like protein (NACSIN), mRNA
 NM_015255 Homo sapiens chromosome 6 open reading frame 133 (C6orf133), mRNA
 NM_015259 Homo sapiens inducible T-cell co-stimulator ligand (ICOSL), mRNA
 NM_015261 Homo sapiens KIAA0056 protein (KIAA0056), mRNA
 NM_015263 Homo sapiens rabconnectin-3 (RC3), mRNA
 NM_015265 Homo sapiens SATB family member 2 (SATB2), mRNA
 NM_015266 Homo sapiens solute carrier family 9 (sodium/hydrogen exchanger), isoform

NM_015268 Homo sapiens DnaJ (Hsp40) homolog, subfamily C, member 13 (DNAJC13),
 NM_015274 Homo sapiens mannosidase, alpha, class 2B, member 2 (MAN2B2), mRNA
 NM_015275 Homo sapiens KIAA1033 protein (KIAA1033), mRNA
 NM_015278 Homo sapiens SAM and SH3 domain containing 1 (SASH1), mRNA
 NM_015281 Homo sapiens KIAA1043 protein (KIAA1043), mRNA
 NM_015282 Homo sapiens cytoplasmic linker associated protein 1 (CLASP1), mRNA
 NM_015284 Homo sapiens KIAA0467 protein (KIAA0467), mRNA
 NM_015286 Homo sapiens desmuslin (DMN), transcript variant B, mRNA
 NM_015289 Homo sapiens vacuolar protein sorting 39 (yeast) (VPS39), mRNA
 NM_015293 Homo sapiens spectrin repeat containing, nuclear envelope 1 (SYNE1), trans
 NM_015296 Homo sapiens dedicator of cytokinesis 9 (DOCK9), mRNA
 NM_015305 Homo sapiens KIAA0759 (KIAA0759), mRNA
 NM_015308 Homo sapiens formin binding protein 4 (FBNP4), mRNA
 NM_015315 Homo sapiens likely ortholog of mouse Ia related protein (LARP), mRNA
 NM_015316 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 13B (PPF
 NM_015319 Homo sapiens tensin like C1 domain containing phosphatase (TENC1), trans
 NM_015321 Homo sapiens mucopidermoid carcinoma translocated 1 (MECT1), mRNA
 NM_015323 Homo sapiens KIAA0776 (KIAA0776), mRNA
 NM_015327 Homo sapiens Est1p-like protein B (EST1B), mRNA
 NM_015328 Homo sapiens KIAA0828 protein (KIAA0828), mRNA
 NM_015329 Homo sapiens KIAA0892 (KIAA0892), mRNA
 NM_015330 Homo sapiens KIAA0376 protein (KIAA0376), mRNA
 NM_015331 Homo sapiens nicastrin (NCSTN), mRNA
 NM_015335 Homo sapiens thyroid hormone receptor associated protein 2 (THRAP2), mF
 NM_015336 Homo sapiens zinc finger, DHHC domain containing 17 (ZDHHC17), mRNA
 NM_015338 Homo sapiens additional sex combs like 1 (Drosophila) (ASXL1), mRNA
 NM_015341 Homo sapiens barren homolog (Drosophila) (BRRN1), mRNA
 NM_015342 Homo sapiens KIAA0073 protein (KIAA0073), mRNA
 NM_015345 Homo sapiens dishevelled associated activator of morphogenesis 2 (DAAM2
 NM_015346 Homo sapiens zinc finger, FYVE domain containing 26 (ZFYE26), mRNA
 NM_015347 Homo sapiens RIM binding protein 2 (KIAA0318), mRNA
 NM_015350 Homo sapiens T-cell activation leucine repeat-rich protein (TA-LRRP), mRNA
 NM_015352 Homo sapiens protein O-fucosyltransferase 1 (POFUT1), transcript variant 1,
 NM_015358 Homo sapiens zinc finger, CW-type with coiled-coil domain 3 (ZCWC3), mF
 NM_015359 Homo sapiens solute carrier family 39 (zinc transporter), member 14 (SLC39
 NM_015360 Homo sapiens KIAA0052 (KIAA0052), mRNA
 NM_015374 Homo sapiens unc-84 homolog B (C. elegans) (UNC84B), mRNA
 NM_015375 Homo sapiens receptor interacting protein kinase 5 (RIPK5), transcript varian
 NM_015378 Homo sapiens vacuolar protein sorting 13D (yeast) (VPS13D), mRNA
 NM_015381 Homo sapiens TAF_A protein 5 (TAF_A5), mRNA
 NM_015382 Homo sapiens HECT domain containing 1 (HECTD1), mRNA
 NM_015386 Homo sapiens component of oligomeric golgi complex 4 (COG4), mRNA
 NM_015391 Homo sapiens anaphase promoting complex subunit 13 (ANAPC13), mRNA
 NM_015395 Homo sapiens DKFZP434B0335 protein (DKFZP434B0335), mRNA
 NM_015397 Homo sapiens WD repeat domain 40A (WDR40A), mRNA
 NM_015404 Homo sapiens deafness, autosomal recessive 31 (DFNB31), mRNA
 NM_015411 Homo sapiens sulfatase modifying factor 2 (SUMF2), mRNA
 NM_015412 Homo sapiens DKFZP434F2021 protein (DKFZP434F2021), mRNA
 NM_015430 Homo sapiens regeneration associated muscle protease (DKFZP586H1213)
 NM_015431 Homo sapiens BIA2 (BIA2), mRNA
 NM_015433 Homo sapiens hepatocellularcarcinoma-associated antigen HCA557a (DKFZ
 NM_015436 Homo sapiens ring finger and CHY zinc finger domain containing 1 (RCHY1)
 NM_015439 Homo sapiens chromosome 6 open reading frame 80 (C6orf80), mRNA
 NM_015440 Homo sapiens formyltetrahydrofolate synthetase domain containing 1 (FTHF
 NM_015441 Homo sapiens olfactomedin-like 2B (OLFM2B), mRNA
 NM_015443 Homo sapiens hypothetical protein LOC284058 (LOC284058), mRNA
 NM_015444 Homo sapiens Ras-induced senescence 1 (RIS1), mRNA

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NM_015446 Homo sapiens ELYS transcription factor-like protein TMBS62 (ELYS), mRNA
 NM_015447 Homo sapiens calmodulin regulated spectrin-associated protein 1 (CAMSAP
 NM_015448 Homo sapiens deleted in a mouse model of primary ciliary dyskinesia (DPC
 NM_015457 Homo sapiens zinc finger, DHHC domain containing 5 (ZDHHC5), mRNA
 NM_015459 Homo sapiens DKFZP564J0863 protein (DKFZP564J0863), mRNA
 NM_015460 Homo sapiens myosin VIIA and Rab interacting protein (MYRIIP), mRNA
 NM_015461 Homo sapiens zinc finger protein 521 (ZNF521), mRNA
 NM_015463 Homo sapiens chromosome 2 open reading frame 32 (C2orf32), mRNA
 NM_015464 Homo sapiens sclerostin domain containing 1 (SOSTDC1), mRNA
 NM_015465 Homo sapiens gem (nuclear organelle) associated protein 5 (GEMIN5), mR
 NM_015466 Homo sapiens protein tyrosine phosphatase, non-receptor type 23 (PTPN23)
 NM_015469 Homo sapiens nipsnap homolog 3A (C. elegans) (NIPSNAP3A), mRNA
 NM_015470 Homo sapiens RAB11 family interacting protein 5 (class I) (RAB11FIP5), mR
 NM_015475 Homo sapiens DKFZP564F0522 protein (DKFZP564F0522), mRNA
 NM_015476 Homo sapiens chromosome 18 open reading frame 10 (C18orf10), mRNA
 NM_015477 Homo sapiens SIN3 homolog A, transcriptional regulator (yeast) (SIN3A), m
 NM_015481 Homo sapiens zinc finger protein 385 (ZNF385), mRNA
 NM_015483 Homo sapiens kelch repeat and BTB (POZ) domain containing 2 (KBTBD2),
 NM_015488 Homo sapiens myofibrillogenesis regulator 1 (MR-1), mRNA
 NM_015503 Homo sapiens SH2-B homolog (SH2B), mRNA
 NM_015508 Homo sapiens TCDD-inducible poly(ADP-ribose) polymerase (TIPARP), mR
 NM_015518 Homo sapiens DKFZP434C131 protein (DKFZP434C131), mRNA
 NM_015522 Homo sapiens dynein 2 light intermediate chain (D2LIC), transcript variant 2,
 NM_015529 Homo sapiens monooxygenase, DBH-like 1 (MOXD1), mRNA
 NM_015531 Homo sapiens DKFZP566P0123 protein (DKFZP566P0123), mRNA
 NM_015532 Homo sapiens glutamate receptor, ionotropic, N-methyl D-aspartate-like 1A (
 NM_015534 Homo sapiens zinc finger, ZZ domain containing 3 (ZZZ3), mRNA
 NM_015541 Homo sapiens leucine-rich repeats and immunoglobulin-like domains 1 (LR
 NM_015547 Homo sapiens thioesterase, adipose associated (THEA), transcript variant 1,
 NM_015548 Homo sapiens dystonin (DST), transcript variant 1eA, mRNA
 NM_015553 Homo sapiens phosphoinositide-binding protein PIP3-E (PIP3-E), mRNA
 NM_015555 Homo sapiens zinc finger protein 451 (ZNF451), mRNA
 NM_015557 Homo sapiens chromodomain helicase DNA binding protein 5 (CHD5), mR
 NM_015558 Homo sapiens synovial sarcoma translocation gene on chromosome 18-like
 NM_015560 Homo sapiens optic atrophy 1 (autosomal dominant) (OPA1), nuclear gene e
 NM_015565 Homo sapiens zinc finger protein 294 (ZNF294), mRNA
 NM_015567 Homo sapiens SLIT and NTRK-like family, member 5 (SLITRK5), mRNA
 NM_015568 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 16B (PP
 NM_015569 Homo sapiens dynamin 3 (DNM3), mRNA
 NM_015575 Homo sapiens trinucleotide repeat containing 15 (TNRC15), mRNA
 NM_015576 Homo sapiens CAZ-associated structural protein (CAST), mRNA
 NM_015578 Homo sapiens chromosome 19 open reading frame 13 (C19orf13), mRNA
 NM_015585 Homo sapiens chromosome 20 open reading frame 26 (C20orf26), mRNA
 NM_015589 Homo sapiens sterile alpha motif domain containing 4 (SAMD4), mRNA
 NM_015597 Homo sapiens G-protein signalling modulator 1 (AGS3-like, C. elegans) (GP
 NM_015600 Homo sapiens chromosome 20 open reading frame 22 (C20orf22), mRNA
 NM_015602 Homo sapiens lamina-associated polypeptide 1B (LAP1B), mRNA
 NM_015604 Homo sapiens WD repeat domain 21 (WDR21), transcript variant 1, mRNA
 NM_015605 Homo sapiens DKFZP566K0524 protein (DKFZP566K0524), mRNA
 NM_015608 Homo sapiens chromosome 10 open reading frame 137 (C10orf137), mRNA
 NM_015609 Homo sapiens putative MAPK activating protein PM20, PM21 (DKFZP566C04
 NM_015617 Homo sapiens pygopus 1 (PYGO1), mRNA
 NM_015627 Homo sapiens LDL receptor adaptor protein (ARH), mRNA
 NM_015631 Homo sapiens chromosome 10 open reading frame 61 (C10orf61), mRNA
 NM_015633 Homo sapiens FGFR1 oncogene partner 2 (FGFR1OP2), mRNA
 NM_015634 Homo sapiens KIAA1279 (KIAA1279), mRNA
 NM_015635 Homo sapiens DKFZP434C212 protein (DKFZP434C212), mRNA

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NM_015639 Homo sapiens GTPase activating RANGAP domain-like 2 pseudogene (GAF
NM_015649 Homo sapiens interferon regulatory factor 2 binding protein 1 (IRF2BP1), mRNA
NM_015655 Homo sapiens zinc finger protein 337 (ZNF337), mRNA
NM_015659 Homo sapiens DKFZP564M182 protein (DKFZP564M182), mRNA
NM_015660 Homo sapiens immunity associated protein 2 (HIMAP2), mRNA
NM_015662 Homo sapiens selective LIM binding factor, rat homolog (SLB), mRNA
NM_015666 Homo sapiens GTP binding protein 5 (putative) (GTPBP5), mRNA
NM_015667 Homo sapiens chromosome 9 open reading frame 36 (C9orf36), mRNA
NM_015668 Homo sapiens DKFZP434I092 protein (DKFZP434I092), mRNA
NM_015687 Homo sapiens filamin A interacting protein 1 (FILIP1), mRNA
NM_015690 Homo sapiens serine/threonine kinase 36 (fused homolog, Drosophila) (STK
NM_015691 Homo sapiens KIAA1280 protein (KIAA1280), mRNA
NM_015692 Homo sapiens C3 and PZP-like, alpha-2-macroglobulin domain containing 8
NM_015693 Homo sapiens PDZ domain containing 6 (PDZK6), mRNA
NM_015694 Homo sapiens KIAA1285 protein (KIAA1285), mRNA
NM_015713 Homo sapiens ribonucleotide reductase M2 B (TP53 inducible) (RRM2B), mRNA
NM_015723 Homo sapiens intracellular membrane-associated calcium-independent phosph
NM_015905 Homo sapiens transcriptional intermediary factor 1 (TIF1), transcript variant 1
NM_015979 Homo sapiens cofactor required for Sp1 transcriptional activation, subunit 3,
NM_016105 Homo sapiens FK506 binding protein 7 (FKBP7), transcript variant 1, mRNA
NM_016133 Homo sapiens insulin induced gene 2 (INSIG2), mRNA
NM_016320 Homo sapiens nucleoporin 98kDa (NUP98), transcript variant 1, mRNA
NM_016544 Homo sapiens Ras-associated protein Rap1 (RBJ), mRNA
NM_017419 Homo sapiens amiloride-sensitive cation channel 5, intestinal (ACCN5), mRNA
NM_017437 Homo sapiens cleavage and polyadenylation specific factor 2, 100kDa (CPSI
NM_017440 Homo sapiens nuclear protein double minute 1 (MDM1), mRNA
NM_017510 Homo sapiens gp25L2 protein (HSGP25L2G), mRNA
NM_017516 Homo sapiens RAB39, member RAS oncogene family (RAB39), mRNA
NM_017519 Homo sapiens AT rich interactive domain 1B (SW11-like) (ARID1B), transcript
NM_017520 Homo sapiens M-phase phosphoprotein, mpp8 (HSMPP8), mRNA
NM_017525 Homo sapiens myotonic dystrophy protein kinase like protein (HSMOPKIN), r
NM_017527 Homo sapiens cDNA for differentially expressed CO16 gene (LY6K), mRNA
NM_017539 Homo sapiens dynein, axonemal, heavy polypeptide 3 (DNAH3), mRNA
NM_017549 Homo sapiens upregulated in colorectal cancer gene 1 (UCC1), mRNA
NM_017550 Homo sapiens KIAA1193 (KIAA1193), mRNA
NM_017553 Homo sapiens homolog of yeast INO80 (INO80), transcript variant 1, mRNA
NM_017554 Homo sapiens KIAA1268 protein (KIAA1268), mRNA
NM_017556 Homo sapiens filamin-binding LIM protein-1 (FBLP-1), mRNA
NM_017563 Homo sapiens Interleukin 17 receptor D (IL17RD), mRNA
NM_017565 Homo sapiens family with sequence similarity 20, member A (FAM20A), mRNA
NM_017570 Homo sapiens 5-oxoprolinase (ATP-hydrolysing) (OPLAH), mRNA
NM_017573 Homo sapiens proprotein convertase subtilisin/kexin type 4 (PCSK4), mRNA
NM_017576 Homo sapiens kinesin family member 27 (KIF27), mRNA
NM_017580 Homo sapiens zinc finger, RAN-binding domain containing 1 (ZRANB1), mRNA
NM_017602 Homo sapiens hypothetical protein DKFZp761A052 (DKFZp761A052), mRNA
NM_017619 Homo sapiens U11/U12 snRNP 65K protein (FLJ25070), mRNA
NM_017628 Homo sapiens hypothetical protein FLJ20032 (FLJ20032), mRNA
NM_017641 Homo sapiens kinesin family member 21A (KIF21A), mRNA
NM_017666 Homo sapiens suppressor of hairy wing homolog 3 (Drosophila) (SUHW3), n
NM_017672 Homo sapiens transient receptor potential cation channel, subfamily M, mem
NM_017725 Homo sapiens hypothetical protein FLJ20249 (FLJ20249), transcript variant :
NM_017747 Homo sapiens ankyrin repeat and KH domain containing 1 (ANKHD1), trans
NM_017754 Homo sapiens chromosome 6 open reading frame 107 (C6orf107), mRNA
NM_017758 Homo sapiens hypothetical protein FLJ20308 (FLJ20308), mRNA
NM_017771 Homo sapiens PX domain containing serine/threonine kinase (PXK), mRNA
NM_017804 Homo sapiens CTF8, chromosome transmission fidelity factor 8 homolog (S
NM_017861 Homo sapiens hypothetical protein FLJ20522 (FLJ20522), mRNA

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NM_017871 Homo sapiens hypothetical protein FLJ20542 (FLJ20542), mRNA
 NM_017879 Homo sapiens hypothetical protein FLJ20557 (FLJ20557), mRNA
 NM_017969 Homo sapiens hypothetical protein FLJ10006 (FLJ10006), mRNA
 NM_017978 Homo sapiens ankyrin repeat and KH domain containing 1 (ANKHD1), trans
 NM_018003 Homo sapiens uveal autoantigen with coiled-coil domains and ankyrin repeat
 NM_018069 Homo sapiens hypothetical protein FLJ10352 (FLJ10352), transcript variant :
 NM_018117 Homo sapiens WD repeat domain 11 (WDR11), mRNA
 NM_018151 Homo sapiens telomere-associated protein RIF1 homolog (Rif1), mRNA
 NM_018177 Homo sapiens Nedd4 binding protein 2 (N4BP2), mRNA
 NM_018193 Homo sapiens hypothetical protein FLJ10719 (FLJ10719), mRNA
 NM_018218 Homo sapiens ubiquitin specific protease 40 (USP40), mRNA
 NM_018237 Homo sapiens cell division cycle and apoptosis regulator 1 (CCAR1), mRNA
 NM_018284 Homo sapiens guanylate binding protein 3 (GBP3), mRNA
 NM_018325 Homo sapiens chromosome 9 open reading frame 72 (C9orf72), transcript vs
 NM_018334 Homo sapiens leucine rich repeat neuronal 3 (LRRN3), mRNA
 NM_018369 Homo sapiens DEP domain containing 1B (DEPDC1B), mRNA
 NM_018392 Homo sapiens hypothetical protein FLJ11331 (FLJ11331), mRNA
 NM_018397 Homo sapiens choline dehydrogenase (CHDH), mRNA
 NM_018405 Homo sapiens hypothetical protein, clone 2746033 (HSA272196), mRNA
 NM_018414 Homo sapiens sialyltransferase 7 (alpha-N-acetylneuraminyl-2,3-beta-galact
 NM_018420 Homo sapiens solute carrier family 22 (organic cation transporter), member 1
 NM_018424 Homo sapiens erythrocyte membrane protein band 4.1 like 4B (EPB41L4B),
 NM_018429 Homo sapiens B double prime 1, subunit of RNA polymerase III transcription
 NM_018462 Homo sapiens chromosome 3 open reading frame 10 (C3orf10), mRNA
 NM_018646 Homo sapiens transient receptor potential cation channel, subfamily V, mem
 NM_018689 Homo sapiens KIAA1199 (KIAA1199), mRNA
 NM_018703 Homo sapiens retinoblastoma binding protein 6 (RBBP6), transcript variant 2
 NM_018704 Homo sapiens hypothetical protein DKFZp547A023 (DKFZp547A023), mRN
 NM_018708 Homo sapiens fem-1 homolog a (C.elegans) (FEM1A), mRNA
 NM_018710 Homo sapiens hypothetical protein DKFZp762O076 (DKFZp762O076), mRN
 NM_018711 Homo sapiens hypothetical protein DKFZp761H039 (DKFZp761H039), mRN
 NM_018712 Homo sapiens hypothetical protein DKFZp547C176 (DKFZp547C176), mRN
 NM_018714 Homo sapiens component of oligomeric golgi complex 1 (COG1), mRNA
 NM_018715 Homo sapiens RCC1-like (TD-60), mRNA
 NM_018717 Homo sapiens mastermind-like 3 (Drosophila) (MAML3), mRNA
 NM_018837 Homo sapiens sulfatase 2 (SULF2), transcript variant 1, mRNA
 NM_018847 Homo sapiens kelch-like 9 (Drosophila) (KLHL9), mRNA
 NM_018881 Homo sapiens DnaJ (Hsp40) homolog, subfamily C, member 10 (DNAJC10),
 NM_018897 Homo sapiens sema domain, seven thrombospondin repeats (type 1 and typ
 NM_018998 Homo sapiens F-box and WD-40 domain protein 5 (FBXW5), transcript varia
 NM_018999 Homo sapiens KIAA1128 protein (KIAA1128), mRNA
 NM_019001 Homo sapiens 5'-3' exoribonuclease 1 (XRN1), mRNA
 NM_019007 Homo sapiens hypothetical protein FLJ20811 (FLJ20811), mRNA
 NM_019010 Homo sapiens keratin 20 (KRT20), mRNA
 NM_019015 Homo sapiens chondroitin sulfate glucuronyltransferase (CSGlcA-T), mRNA
 NM_019022 Homo sapiens FLJ20793 protein (FLJ20793), mRNA
 NM_019026 Homo sapiens putative membrane protein (LOC54499), mRNA
 NM_019029 Homo sapiens carboxypeptidase, vitellogenic-like (CPVL), transcript variant :
 NM_019030 Homo sapiens DEAH (Asp-Glu-Ala-His) box polypeptide 29 (DHX29), mRNA
 NM_019032 Homo sapiens thrombospondin repeat containing 1 (TSRC1), mRNA
 NM_019036 Homo sapiens 3-hydroxymethyl-3-methylglutaryl-Coenzyme A lyase-like 1 (H
 NM_019051 Homo sapiens mitochondrial ribosomal protein L50 (MRPL50), nuclear gene
 NM_019053 Homo sapiens SEC15-like 1 (S. cerevisiae) (SEC15L1), mRNA
 NM_019055 Homo sapiens roundabout homolog 4, magic roundabout (Drosophila) (ROB)
 NM_019065 Homo sapiens EF hand calcium binding protein 2 (EFGBP2), mRNA
 NM_019072 Homo sapiens small glutamine-rich tetratricopeptide repeat (TPR)-containing
 NM_019075 Homo sapiens UDP glycosyltransferase 1 family, polypeptide A10 (UGT1A1C

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NM_019077 Homo sapiens UDP glycosyltransferase 1 family, polypeptide A7 (UGT1A7),
 NM_019078 Homo sapiens UDP glycosyltransferase 1 family, polypeptide A5 (UGT1A5),
 NM_019085 Homo sapiens F-box and leucine-rich repeat protein 19 (FBXL19), mRNA
 NM_019092 Homo sapiens hypothetical protein KIAA1164 (KIAA1164), mRNA
 NM_019104 Homo sapiens protein F25965 (F25965), mRNA
 NM_019107 Homo sapiens chromosome 19 open reading frame 10 (C19orf10), mRNA
 NM_019590 Homo sapiens KIAA1217 (KIAA1217), mRNA
 NM_019593 Homo sapiens hypothetical protein KIAA1434 (KIAA1434), mRNA
 NM_019594 Homo sapiens leucine rich repeat containing 8 (LRRCC8), mRNA
 NM_019850 Homo sapiens neuronal guanine nucleotide exchange factor (NGEF), mRNA
 NM_020063 Homo sapiens BarH-like 2 (Drosophila) (BARHL2), mRNA
 NM_020116 Homo sapiens follistatin-like 5 (FSTL5), mRNA
 NM_020124 Homo sapiens interferon, kappa (IFNK), mRNA
 NM_020170 Homo sapiens nicalin (LOC56926), mRNA
 NM_020172 Homo sapiens SPPL2b (SPPL2B), mRNA
 NM_020175 Homo sapiens hypothetical protein from EUROMIMAGE 1967720 (LOC56931)
 NM_020192 Homo sapiens chromosome 7 open reading frame 36 (C7orf36), mRNA
 NM_020204 Homo sapiens LIM homeobox 9 (LHX9), mRNA
 NM_020207 Homo sapiens chromosome 9 open reading frame 102 (C9orf102), mRNA
 NM_020209 Homo sapiens src homology 2 domain-containing transforming protein D (Sf)
 NM_020210 Homo sapiens sema domain, immunoglobulin domain (Ig), transmembrane c
 NM_020211 Homo sapiens RGM domain family, member A (RGMA), mRNA
 NM_020212 Homo sapiens hypothetical protein from EUROMIMAGE 384293 (LOC56964),
 NM_020214 Homo sapiens hypothetical protein from EUROMIMAGE 1977056 (LOC56965)
 NM_020219 Homo sapiens carcinoembryonic antigen-like 1 (CEAL1), mRNA
 NM_020223 Homo sapiens family with sequence similarity 20, member C (FAM20C), mRi
 NM_020311 Homo sapiens chemokine orphan receptor 1 (CMKOR1), mRNA
 NM_020312 Homo sapiens hypothetical protein DKFZp434K046 (DKFZP434K046), mRN
 NM_020318 Homo sapiens pappalysin 2 (PAPPA2), transcript variant 1, mRNA
 NM_020319 Homo sapiens ankyrin repeat and MYND domain containing 2 (ANKMY2), mI
 NM_020320 Homo sapiens arginyl-tRNA synthetase-like (RARS), mRNA
 NM_020336 Homo sapiens KIAA1219 protein (KIAA1219), mRNA
 NM_020338 Homo sapiens retinoic acid induced 17 (RAI17), mRNA
 NM_020340 Homo sapiens KIAA1244 (KIAA1244), mRNA
 NM_020341 Homo sapiens p21(CDKN1A)-activated kinase 7 (PAK7), transcript variant 1,
 NM_020376 Homo sapiens transport-secretion protein 2.2 (TTS-2.2), mRNA
 NM_020378 Homo sapiens K562 cell-derived leucine-zipper-like protein 1 (KLP1), mRNA
 NM_020383 Homo sapiens X-prolyl aminopeptidase (aminopeptidase P) 1, soluble (XPNF
 NM_020409 Homo sapiens mitochondrial ribosomal protein L47 (MRPL47), nuclear gene
 NM_020416 Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit B (PF
 NM_020417 Homo sapiens T-box 20 (TBX20), mRNA
 NM_020420 Homo sapiens deleted in azoospermia 4 (DAZ4), mRNA
 NM_020429 Homo sapiens SMAD specific E3 ubiquitin protein ligase 1 (SMURF1), transc
 NM_020432 Homo sapiens putative homeodomain transcription factor 2 (PHTF2), mRNA
 NM_020436 Homo sapiens dolichyl pyrophosphate phosphatase 1 (DOLPP1), mRNA
 NM_020440 Homo sapiens prostaglandin F2 receptor negative regulator (PTGFRN), mRr
 NM_020447 Homo sapiens chromosome 15 open reading frame 17 (C15orf17), mRNA
 NM_020451 Homo sapiens selenoprotein N, 1 (SEPN1), transcript variant 1, mRNA
 NM_020452 Homo sapiens ATPase, Class I, type 8B, member 2 (ATP8B2), mRNA
 NM_020453 Homo sapiens ATPase, Class V, type 10D (ATP10D), mRNA
 NM_020455 Homo sapiens G protein-coupled receptor 126 (GPR126), mRNA
 NM_020456 Homo sapiens chromosome 13 open reading frame 1 (C13orf1), mRNA
 NM_020457 Homo sapiens THAP domain containing 11 (THAP11), mRNA
 NM_020462 Homo sapiens KIAA1181 protein (KIAA1181), mRNA
 NM_020463 Homo sapiens KIAA1387 protein (KIAA1387), mRNA
 NM_020468 Homo sapiens sorting nexin 14 (SNX14), transcript variant 2, mRNA
 NM_020531 Homo sapiens chromosome 20 open reading frame 3 (C20orf3), mRNA

NM_020532 Homo sapiens reticulon 4 (RTN4), transcript variant 1, mRNA
 NM_020536 Homo sapiens CSRP2 binding protein (CSRP2BP), transcript variant 1, mRNA
 NM_020546 Homo sapiens adenylate cyclase 2 (brain) (ADCY2), mRNA
 NM_020631 Homo sapiens putative NFkB activating protein (KIAA0720), transcript variant
 NM_020647 Homo sapiens junctophilin 1 (JPH1), mRNA
 NM_020693 Homo sapiens Down syndrome cell adhesion molecule like 1 (DSCAML1), mRNA
 NM_020695 Homo sapiens transcription elongation factor B polypeptide 3 binding protein
 NM_020696 Homo sapiens KIAA1143 protein (KIAA1143), mRNA
 NM_020697 Homo sapiens potassium voltage-gated channel, delayed-rectifier, subfamily
 NM_020698 Homo sapiens KIAA1145 protein (KIAA1145), mRNA
 NM_020699 Homo sapiens transcription repressor p66 beta component of the MeCP1 complex
 NM_020701 Homo sapiens KIAA1160 protein (KIAA1160), mRNA
 NM_020702 Homo sapiens KIAA1161 (KIAA1161), mRNA
 NM_020706 Homo sapiens splicing factor, arginine/serine-rich 15 (SFRS15), mRNA
 NM_020710 Homo sapiens KIAA1185 protein (KIAA1185), mRNA
 NM_020713 Homo sapiens KIAA1196 protein (KIAA1196), mRNA
 NM_020714 Homo sapiens zinc finger protein 490 (ZNF490), mRNA
 NM_020718 Homo sapiens ubiquitin specific protease 31 (USP31), mRNA
 NM_020728 Homo sapiens KIAA1228 protein (KIAA1228), mRNA
 NM_020732 Homo sapiens AT rich interactive domain 1B (SWI1-like) (ARID1B), transcript
 NM_020739 Homo sapiens cell cycle progression 1 (CCPG1), mRNA
 NM_020740 Homo sapiens ankyrin repeat and FYVE domain containing 1 (ANKFY1), transcript
 NM_020742 Homo sapiens neuroligin 4, X-linked (NLGN4X), transcript variant 1, mRNA
 NM_020744 Homo sapiens metastasis associated family, member 3 (MTA3), mRNA
 NM_020745 Homo sapiens alanyl-tRNA synthetase like (AARSL), mRNA
 NM_020746 Homo sapiens KIAA1271 protein (KIAA1271), mRNA
 NM_020748 Homo sapiens KIAA1287 protein (KIAA1287), mRNA
 NM_020750 Homo sapiens exportin 5 (XPO5), mRNA
 NM_020751 Homo sapiens component of oligomeric golgi complex 6 (COG6), mRNA
 NM_020752 Homo sapiens G protein-coupled receptor 158 (GPR158), mRNA
 NM_020753 Homo sapiens CASK Interacting protein 2 (CASKIN2), mRNA
 NM_020755 Homo sapiens tumor differentially expressed 2 (TDE2), mRNA
 NM_020761 Homo sapiens raptor (raptor), mRNA
 NM_020762 Homo sapiens SLIT-ROBO Rho GTPase activating protein 1 (SRGAP1), mRNA
 NM_020765 Homo sapiens retinoblastoma-associated factor 600 (RBAF600), mRNA
 NM_020769 Homo sapiens KIAA1318 protein (KIAA1318), mRNA
 NM_020771 Homo sapiens HECT domain and ankyrin repeat containing, E3 ubiquitin protein
 NM_020772 Homo sapiens 82-kD FMRP Interacting Protein (182-FIP), mRNA
 NM_020773 Homo sapiens TBC1 domain family, member 14 (TBC1D14), mRNA
 NM_020774 Homo sapiens mindbomb homolog 1 (Drosophila) (MIB1), mRNA
 NM_020775 Homo sapiens mab1 (KIAA1324), mRNA
 NM_020778 Homo sapiens likely ortholog of mouse myocytic induction/differentiation orig
 NM_020779 Homo sapiens WD repeat domain 35 (WDR35), mRNA
 NM_020781 Homo sapiens zinc finger protein 398 (ZNF398), transcript variant 2, mRNA
 NM_020783 Homo sapiens synaptotagmin IV (SYT4), mRNA
 NM_020786 Homo sapiens pyruvate dehydrogenase phosphatase isoenzyme 2 (PDP2), mRNA
 NM_020787 Homo sapiens zinc finger protein 624 (ZNF624), mRNA
 NM_020789 Homo sapiens immunoglobulin superfamily, member 9 (IGSF9), mRNA
 NM_020791 Homo sapiens serine/threonine protein kinase TAO1 homolog (KIAA1361), mRNA
 NM_020792 Homo sapiens KIAA1363 protein (KIAA1363), mRNA
 NM_020795 Homo sapiens neuroligin 2 (NLGN2), mRNA
 NM_020799 Homo sapiens associated molecule with the SH3 domain of STAM (AMSH) li
 NM_020800 Homo sapiens KIAA1374 protein (KIAA1374), mRNA
 NM_020801 Homo sapiens arrestin domain containing 3 (ARRDC3), mRNA
 NM_020803 Homo sapiens kelch-like 8 (Drosophila) (KLHL8), mRNA
 NM_020804 Homo sapiens protein kinase C and casein kinase substrate in neurons 1 (P
 NM_020808 Homo sapiens signal-induced proliferation-associated 1 like 2 (SIPA1L2), mRNA

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NM_020809 Homo sapiens Rho GTPase activating protein 20 (ARHGAP20), mRNA
 NM_020810 Homo sapiens KIAA1393 (KIAA1393), mRNA
 NM_020812 Homo sapiens dedicator of cytokinesis 6 (DOCK6), mRNA
 NM_020813 Homo sapiens zinc finger protein 471 (ZNF471), mRNA
 NM_020816 Homo sapiens kinesin family member 17 (KIF17), mRNA
 NM_020817 Homo sapiens KIAA1407 protein (KIAA1407), mRNA
 NM_020818 Homo sapiens KIAA1409 (KIAA1409), mRNA
 NM_020820 Homo sapiens phosphatidylinositol 3,4,5-trisphosphate-dependent RAC exct
 NM_020824 Homo sapiens Rho GTPase activating protein 21 (ARHGAP21), mRNA
 NM_020825 Homo sapiens Crm, cramped-like (Drosophila) (CRAMP1L), mRNA
 NM_020826 Homo sapiens synaptotagmin XIII (SYT13), mRNA
 NM_020828 Homo sapiens zinc finger protein 28 homolog (mouse) (ZFP28), mRNA
 NM_020832 Homo sapiens KIAA1441 protein (KIAA1441), mRNA
 NM_020834 Homo sapiens KIAA1443 (KIAA1443), mRNA
 NM_020839 Homo sapiens WD repeat endosomal protein (KIAA1449), mRNA
 NM_020844 Homo sapiens KIAA1456 protein (KIAA1456), mRNA
 NM_020845 Homo sapiens phosphatidylinositol transfer protein, membrane-associated 2
 NM_020847 Homo sapiens trinucleotide repeat containing 6 (TNRC6), mRNA
 NM_020850 Homo sapiens Ran-binding protein 10 (RANBP10), mRNA
 NM_020851 Homo sapiens KIAA1465 protein (KIAA1465), mRNA
 NM_020854 Homo sapiens KIAA1468 (KIAA1468), mRNA
 NM_020856 Homo sapiens zinc finger protein 537 (ZNF537), mRNA
 NM_020858 Homo sapiens sema domain, transmembrane domain (TM), and cytoplasmic
 NM_020859 Homo sapiens Shroom-related protein (ShrmL), mRNA
 NM_020860 Homo sapiens stromal interaction molecule 2 (STIM2), mRNA
 NM_020861 Homo sapiens zinc finger and BTB domain containing 2 (ZBTB2), mRNA
 NM_020863 Homo sapiens zinc finger protein 406 (ZNF406), mRNA
 NM_020867 Homo sapiens ubiquitin associated protein 2 (UBAP2), transcript variant 2, r
 NM_020868 Homo sapiens dipeptidylpeptidase 10 (DPP10), mRNA
 NM_020870 Homo sapiens SH3 multiple domains 2 (SH3MD2), mRNA
 NM_020871 Homo sapiens leucine-rich repeats and calponin homology (CH) domain con
 NM_020873 Homo sapiens leucine rich repeat neuronal 1 (LRRN1), mRNA
 NM_020875 Homo sapiens Fraser syndrome 1 (FRAS1), transcript variant 3, mRNA
 NM_020880 Homo sapiens zinc finger protein 530 (ZNF530), mRNA
 NM_020882 Homo sapiens KIAA1510 protein (KIAA1510), mRNA
 NM_020889 Homo sapiens PHD finger protein 12 (PHF12), mRNA
 NM_020890 Homo sapiens KIAA1524 protein (KIAA1524), mRNA
 NM_020892 Homo sapiens deltex homolog 2 (Drosophila) (DTX2), mRNA
 NM_020895 Homo sapiens KIAA1533 (KIAA1533), mRNA
 NM_020897 Homo sapiens hyperpolarization activated cyclic nucleotide-gated potassium
 NM_020899 Homo sapiens zinc finger and BTB domain containing 4 (ZBTB4), mRNA
 NM_020914 Homo sapiens chromosome 17 open reading frame 27 (C17orf27), mRNA
 NM_020918 Homo sapiens glycerol-3-phosphate acyltransferase, mitochondrial (GPAM),
 NM_020922 Homo sapiens protein kinase, lysine deficient 3 (PRKWNK3), transcript varia
 NM_020925 Homo sapiens KIAA1573 protein (KIAA1573), mRNA
 NM_020926 Homo sapiens BCL6 co-repressor (BCOR), transcript variant 2, mRNA
 NM_020927 Homo sapiens KIAA1576 protein (KIAA1576), mRNA
 NM_020932 Homo sapiens melanoma antigen, family E, 1 (MAGEE1), mRNA
 NM_020935 Homo sapiens ubiquitin specific protease 37 (USP37), mRNA
 NM_020936 Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 55 (DDX55), mRNA
 NM_020939 Homo sapiens copine V (CPN5E), mRNA
 NM_020944 Homo sapiens glucosidase, beta (bile acid) 2 (GBA2), mRNA
 NM_020947 Homo sapiens KIAA1609 protein (KIAA1609), mRNA
 NM_020948 Homo sapiens mesoderm induction early response 1 (MI-ER1), mRNA
 NM_020951 Homo sapiens zinc finger protein 529 (ZNF529), mRNA
 NM_020952 Homo sapiens transient receptor potential cation channel, subfamily M, mem
 NM_020954 Homo sapiens KIAA1618 (KIAA1618), mRNA

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NM_020961 Homo sapiens KIAA1627 protein (KIAA1627), mRNA
 NM_020962 Homo sapiens likely ortholog of mouse neighbor of Punc E11 (NOPE), mRNA
 NM_020964 Homo sapiens KIAA1632 protein (KIAA1632), mRNA
 NM_020965 Homo sapiens membrane-associated guanylate kinase-related (MAGI-3) (M)
 NM_020970 Homo sapiens KIAA1641 (KIAA1641), mRNA
 NM_020971 Homo sapiens spectrin, beta, non-erythrocytic 4 (SPTBN4), mRNA
 NM_021006 Homo sapiens chemokine (C-C motif) ligand 3-like 1 (CCL3L1), mRNA
 NM_021009 Homo sapiens ubiquitin C (UBC), mRNA
 NM_021035 Homo sapiens KIAA1404 protein (KIAA1404), mRNA
 NM_021044 Homo sapiens desert hedgehog homolog (Drosophila) (DHH), mRNA
 NM_021045 Homo sapiens zinc finger protein 248 (ZNF248), mRNA
 NM_021059 Homo sapiens histone 2, H3c (HIST2H3C), mRNA
 NM_021061 Homo sapiens zinc finger protein 647 (ZNF647), mRNA
 NM_021072 Homo sapiens hyperpolarization activated cyclic nucleotide-gated potassium
 NM_021088 Homo sapiens zinc finger protein 2 (A1-5) (ZNF2), mRNA
 NM_021089 Homo sapiens zinc finger protein 8 (clone HF.18) (ZNF8), mRNA
 NM_021116 Homo sapiens adenylate cyclase 1 (brain) (ADCY1), mRNA
 NM_021117 Homo sapiens cryptochrome 2 (photolyase-like) (CRY2), mRNA
 NM_021143 Homo sapiens zinc finger protein 20 (KOX 13) (ZNF20), mRNA
 NM_021148 Homo sapiens zinc finger protein 273 (ZNF273), mRNA
 NM_021149 Homo sapiens coactosin-like 1 (Dictyostelium) (COTL1), mRNA
 NM_021184 Homo sapiens splicing factor 4 (SF4), transcript variant b, mRNA
 NM_021185 Homo sapiens hypothetical protein from clone 24828 (KIAA1747), mRNA
 NM_021180 Homo sapiens transcription factor CP2-like 4 (TFCP2L4), transcript variant 1
 NM_021202 Homo sapiens tumor protein p53 inducible nuclear protein 2 (TP53INP2), mF
 NM_021217 Homo sapiens zinc finger protein 77 (pT1) (ZNF77), mRNA
 NM_021218 Homo sapiens chromosome 9 open reading frame 80 (C9orf80), mRNA
 NM_021222 Homo sapiens Tcd37 homolog (HTCD37), mRNA
 NM_021224 Homo sapiens zinc finger protein 462 (ZNF462), mRNA
 NM_021227 Homo sapiens DC2 protein (DC2), mRNA
 NM_021228 Homo sapiens serine arginine-rich pre-mRNA splicing factor SR-A1 (SR-A1),
 NM_021237 Homo sapiens selenoprotein K (SELK), mRNA
 NM_021250 Homo sapiens leukocyte Ig-like receptor 9 (LIR9), transcript variant 1, mRNA
 NM_021260 Homo sapiens zinc finger, FYVE domain containing 1 (ZFYE1), transcript v
 NM_021636 Homo sapiens leucine-rich repeat-containing G protein-coupled receptor 6 L
 NM_021649 Homo sapiens toll-like receptor adaptor molecule 2 (TICAM2), mRNA
 NM_021652 Homo sapiens SMA4 (SMA4), mRNA
 NM_021915 Homo sapiens zinc finger protein 69 (Cos5) (ZNF69), mRNA
 NM_021916 Homo sapiens zinc finger protein 70 (Cos17) (ZNF70), mRNA
 NM_021936 Homo sapiens pappalysin 2 (PAPPA2), transcript variant 2, mRNA
 NM_021937 Homo sapiens elongation factor for selenoprotein translation (SELB), mRNA
 NM_022045 Homo sapiens Mdm2, transformed 3T3 cell double minute 2, p53 binding pro
 NM_022075 Homo sapiens LAG1 longevity assurance homolog 2 (S. cerevisiae) (LASS2)
 NM_022080 Homo sapiens N-ethylmaleimide-sensitive factor attachment protein, beta (N
 NM_022085 Homo sapiens thioredoxin domain containing 5 (TXNDC5), transcript variant
 NM_022092 Homo sapiens CTF18, chromosome transmission fidelity factor 18 homolog
 NM_022106 Homo sapiens chromosome 20 open reading frame 177 (C20orf177), mRNA
 NM_022115 Homo sapiens PR domain containing 15 (PRDM15), mRNA
 NM_022138 Homo sapiens SPARC related modular calcium binding 2 (SMOC2), mRNA
 NM_022160 Homo sapiens DMRT-like family A1 (DMRTA1), mRNA
 NM_022166 Homo sapiens xylosyltransferase 1 (XYLT1), mRNA
 NM_022351 Homo sapiens EF hand calcium binding protein 1 (EFCBP1), mRNA
 NM_022475 Homo sapiens hedgehog interacting protein (HHIP), mRNA
 NM_022478 Homo sapiens cadherin-like 24 (CDH24), mRNA
 NM_022479 Homo sapiens Williams-Beuren syndrome chromosome region 17 (WBSCR1
 NM_022486 Homo sapiens sushi domain containing 1 (SUSD1), mRNA
 NM_022491 Homo sapiens likely ortholog of mouse Sds3 (SDS3), mRNA

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NM_022572 Homo sapiens myofibrillogenesis regulator 1 (MR-1), mRNA
 NM_022733 Homo sapiens hypothetical protein AL133206 (LOC84744), mRNA
 NM_022742 Homo sapiens hypothetical protein DKFZP434G156 (DKFZP434G156), mRNA
 NM_022745 Homo sapiens ATP synthase mitochondrial F1 complex assembly factor 1 (A
 NM_022757 Homo sapiens hypothetical protein FLJ12892 (FLJ12892), mRNA
 NM_022824 Homo sapiens F-box and leucine-rich repeat protein 17 (FBXL17), mRNA
 NM_022833 Homo sapiens chromosome 9 open reading frame 88 (C9orf88), mRNA
 NM_022835 Homo sapiens likely ortholog of mouse common-site lymphoma/leukemia GE
 NM_022913 Homo sapiens vasculin (DKFZp761C169), mRNA
 NM_023002 Homo sapiens hyaluronan and proteoglycan link protein 4 (HAPLN4), mRNA
 NM_023006 Homo sapiens kallikrein 15 (KLK15), transcript variant 1, mRNA
 NM_023939 Homo sapiens hypothetical protein MGC2752 (MGC2752), mRNA
 NM_023943 Homo sapiens hypothetical protein MGC3040 (MGC3040), mRNA
 NM_024007 Homo sapiens early B-cell factor (EBF), mRNA
 NM_024019 Homo sapiens neurogenin 2 (NEUROG2), mRNA
 NM_024100 Homo sapiens WD repeat domain 18 (WDR18), mRNA
 NM_024316 Homo sapiens leukocyte receptor cluster (LRC) member 1 (LENG1), mRNA
 NM_024335 Homo sapiens iroquois homeobox protein 6 (IRX6), mRNA
 NM_024336 Homo sapiens iroquois homeobox protein 3 (IRX3), mRNA
 NM_024342 Homo sapiens glucocorticoid receptor DNA binding factor 1 (GRLF1), transcr
 NM_024344 Homo sapiens calpain 3, (p94) (CAPN3), transcript variant 2, mRNA
 NM_024420 Homo sapiens phospholipase A2, group IVA (cytosolic, calcium-dependent) i
 NM_024493 Homo sapiens zinc finger protein 306 (ZNF306), mRNA
 NM_024496 Homo sapiens chromosome 14 open reading frame 4 (C14orf4), mRNA
 NM_024511 Homo sapiens chromosome 4 open reading frame 15 (C4orf15), mRNA
 NM_024517 Homo sapiens PHD finger protein 2 (PHF2), transcript variant 2, mRNA
 NM_024563 Homo sapiens hypothetical protein FLJ20097 (FLJ20097), mRNA
 NM_024621 Homo sapiens hypothetical protein FLJ12604 (FLJ12604), mRNA
 NM_024625 Homo sapiens zinc finger CCHC type, antiviral 1 (ZC3HAV1), transcript varia
 NM_024684 Homo sapiens PTD015 protein (PTD015), mRNA
 NM_024742 Homo sapiens armadillo repeat containing 5 (ARMCS), mRNA
 NM_024769 Homo sapiens adipocyte-specific adhesion molecule (ASAM), mRNA
 NM_024870 Homo sapiens DEP domain containing 2 (DEPDC2), transcript variant 1, mR
 NM_024878 Homo sapiens CGI-72 protein (CGI-72), transcript variant 4, mRNA
 NM_024933 Homo sapiens hypothetical protein FLJ12056 (FLJ12056), mRNA
 NM_024953 Homo sapiens hypothetical protein FLJ13089 (FLJ13089), mRNA
 NM_025169 Homo sapiens zinc finger protein 167 (ZNF167), transcript variant 2, mRNA
 NM_025196 Homo sapiens GrpE-like 1, mitochondrial (E. coli) (GRPEL1), mRNA
 NM_025202 Homo sapiens EF hand domain containing 1 (EFHD1), mRNA
 NM_025219 Homo sapiens DnaJ (Hsp40) homolog, subfamily C, member 5 (DNAJC5), m
 NM_025224 Homo sapiens BTB (POZ) domain containing 4 (BTBD4), mRNA
 NM_025248 Homo sapiens SNAP25-interacting protein (SNIP), mRNA
 NM_025252 Homo sapiens Ras association (RalGDS/AF-6) and pleckstrin homology dom
 NM_025256 Homo sapiens HLA-B associated transcript 8 (BAT8), transcript variant NG3
 NM_030625 Homo sapiens CXXC finger 6 (CXXC6), mRNA
 NM_030627 Homo sapiens cytoplasmic polyadenylation element binding protein 4 (CPEB
 NM_030628 Homo sapiens KIAA1698 protein (KIAA1698), mRNA
 NM_030629 Homo sapiens c-Maf-inducing protein (CMIP), transcript variant Tc-mip, mR
 NM_030630 Homo sapiens chromosome 17 open reading frame 28 (C17orf28), mRNA
 NM_030633 Homo sapiens KIAA1712 (KIAA1712), mRNA
 NM_030634 Homo sapiens zinc finger protein 436 (ZNF436), mRNA
 NM_030636 Homo sapiens KIAA1706 protein (KIAA1706), mRNA
 NM_030637 Homo sapiens DDHD domain containing 1 (DDHD1), mRNA
 NM_030639 Homo sapiens KIAA1701 protein (KIAA1701), mRNA
 NM_030640 Homo sapiens dual specificity phosphatase 16 (DUSP16), mRNA
 NM_030644 Homo sapiens apolipoprotein L, 3 (APOL3), transcript variant alpha/b, mRNA
 NM_030645 Homo sapiens KIAA1720 protein (KIAA1720), mRNA

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NM_030650 Homo sapiens KIAA1715 (KIAA1715), mRNA
 NM_030789 Homo sapiens histocompatibility (minor) 13 (HM13), transcript variant 1, mRNA
 NM_030812 Homo sapiens actin like protein (LOC81569), mRNA
 NM_030883 Homo sapiens olfactory receptor, family 2, subfamily H, member 1 (OR2H1), mRNA
 NM_030906 Homo sapiens serine/threonine kinase 33 (STK33), mRNA
 NM_030919 Homo sapiens chromosome 20 open reading frame 129 (C20orf129), mRNA
 NM_030922 Homo sapiens non-imprinted in Prader-Willi/Angelman syndrome 2 (NIPAD2), mRNA
 NM_030923 Homo sapiens hypothetical protein DKFZp566N034 (DKFZP566N034), mRNA
 NM_030949 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 14C (PP1), mRNA
 NM_030957 Homo sapiens a disintegrin-like and metalloprotease (repolyrin type) with th, mRNA
 NM_030961 Homo sapiens tripartite motif-containing 56 (TRIM56), mRNA
 NM_030962 Homo sapiens Charcot-Marie-Tooth neuropathy 4B2 (autosomal recessive), v, mRNA
 NM_031303 Homo sapiens similar to RIKEN cDNA 4933439B08 gene (MGC33211), mRNA
 NM_031444 Homo sapiens chromosome 22 open reading frame 13 (C22orf13), mRNA
 NM_031448 Homo sapiens chromosome 19 open reading frame 12 (C19orf12), mRNA
 NM_031454 Homo sapiens selenoprotein O (SELO), mRNA
 NM_031467 Homo sapiens solute carrier family 4, sodium bicarbonate cotransporter, mer, mRNA
 NM_031490 Homo sapiens peroxisomal lon protease (LONP), mRNA
 NM_031888 Homo sapiens pro-melanin-concentrating hormone-like 2 (PMCHL2), mRNA
 NM_031895 Homo sapiens calcium channel, voltage-dependent, gamma subunit 8 (CAC), mRNA
 NM_031912 Homo sapiens synaptotagmin XV (SYT15), transcript variant a, mRNA
 NM_031913 Homo sapiens chr3 synaptotagmin (CHR3SYT), mRNA
 NM_031914 Homo sapiens synaptotagmin XIV-like (SYT14L), mRNA
 NM_031935 Homo sapiens hemicentin (FIBL-6), mRNA
 NM_032017 Homo sapiens Ser/Thr-like kinase (MGC4796), mRNA
 NM_032111 Homo sapiens mitochondrial ribosomal protein L14 (MRPL14), nuclear gene, mRNA
 NM_032119 Homo sapiens monogenic, audiogenic seizure susceptibility 1 homolog (mou), mRNA
 NM_032123 Homo sapiens kln of IRRE like 2 (Drosophila) (KIRREL2), transcript variant 1, mRNA
 NM_032132 Homo sapiens HORMA domain containing protein (NOHMA), mRNA
 NM_032137 Homo sapiens hypothetical protein DKFZp434N1817 (DKFZP434N1817), mRNA
 NM_032156 Homo sapiens C1q domain containing 1 (C1QDC1), transcript variant 3, mRNA
 NM_032160 Homo sapiens chromosome 18 open reading frame 4 (C18orf4), mRNA
 NM_032165 Homo sapiens hypothetical protein FLJ12303 (FLJ12303), mRNA
 NM_032168 Homo sapiens hypothetical protein FLJ12519 (FLJ12519), mRNA
 NM_032194 Homo sapiens brx domain containing 1 (BXDC1), mRNA
 NM_032195 Homo sapiens SON DNA binding protein (SON), transcript variant b, mRNA
 NM_032217 Homo sapiens ankyrin repeat domain 17 (ANKRD17), transcript variant 1, mRNA
 NM_032222 Homo sapiens hypothetical protein FLJ22374 (FLJ22374), mRNA
 NM_032226 Homo sapiens zinc finger, CCHC domain containing 7 (ZCCHC7), mRNA
 NM_032228 Homo sapiens male sterility domain containing 2 (MLSTD2), mRNA
 NM_032230 Homo sapiens hypothetical protein FLJ22789 (FLJ22789), mRNA
 NM_032279 Homo sapiens hypothetical protein DKFZp761I1011 (DKFZp761I1011), mRNA
 NM_032282 Homo sapiens hypothetical protein DKFZp547D155 (DKFZp547D155), mRNA
 NM_032283 Homo sapiens zinc finger, DHHC domain containing 18 (ZDHHC18), mRNA
 NM_032285 Homo sapiens hypothetical protein MGC3207 (MGC3207), mRNA
 NM_032286 Homo sapiens hypothetical protein MGC5309 (MGC5309), mRNA
 NM_032422 Homo sapiens G protein-coupled receptor 123 (GPR123), mRNA
 NM_032423 Homo sapiens zinc finger protein 528 (ZNF528), mRNA
 NM_032425 Homo sapiens KIAA1822 (KIAA1822), mRNA
 NM_032427 Homo sapiens mastermind-like 2 (Drosophila) (MAML2), mRNA
 NM_032429 Homo sapiens leucine zipper, putative tumor suppressor 2 (LZTS2), mRNA
 NM_032430 Homo sapiens KIAA1811 protein (KIAA1811), mRNA
 NM_032431 Homo sapiens HRD1 protein (HRD1), transcript variant 1, mRNA
 NM_032432 Homo sapiens actin binding LIM protein family, member 2 (ABLIM2), mRNA
 NM_032433 Homo sapiens zinc finger protein 333 (ZNF333), mRNA
 NM_032434 Homo sapiens zinc finger protein 512 (ZNF512), mRNA
 NM_032435 Homo sapiens mixed lineage kinase 4 (KIAA1804), mRNA

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NM_032436 Homo sapiens chromosome 13 open reading frame 8 (C13orf8), mRNA
 NM_032439 Homo sapiens phytanoyl-CoA hydroxylase interacting protein-like (PHYHPL), mRNA
 NM_032440 Homo sapiens ligand-dependent corepressor (MLR2), mRNA
 NM_032444 Homo sapiens BTB (POZ) domain containing 12 (BTBD12), mRNA
 NM_032448 Homo sapiens KIAA1838 (KIAA1838), mRNA
 NM_032452 Homo sapiens junctophilin 4 (JPH4), mRNA
 NM_032458 Homo sapiens PHD finger protein 6 (PHF6), mRNA
 NM_032477 Homo sapiens mitochondrial ribosomal protein L41 (MRPL41), nuclear gene
 NM_032478 Homo sapiens mitochondrial ribosomal protein L38 (MRPL38), nuclear gene
 NM_032479 Homo sapiens mitochondrial ribosomal protein L36 (MRPL36), nuclear gene
 NM_032482 Homo sapiens DOT1-like, histone H3 methyltransferase (S. cerevisiae) (DOT)
 NM_032497 Homo sapiens zinc finger protein 559 (ZNF559), mRNA
 NM_032501 Homo sapiens acetyl-Coenzyme A synthetase 2 (AMP forming)-like (ACAS2)
 NM_032505 Homo sapiens T-cell activation kelch repeat protein (TA-KRP), mRNA
 NM_032506 Homo sapiens KIAA1841 protein (KIAA1841), mRNA
 NM_032508 Homo sapiens family with sequence similarity 11, member A (FAM11A), mR
 NM_032511 Homo sapiens chromosome 6 open reading frame 168 (C6orf168), mRNA
 NM_032512 Homo sapiens PDZ domain containing 4 (PDZK4), mRNA
 NM_032517 Homo sapiens lysozyme-like 1 (LYZL1), mRNA
 NM_032528 Homo sapiens beta-galactoside alpha-2,6-sialyltransferase II (ST6GalII), mR
 NM_032531 Homo sapiens kin of IRRE like 3 (Drosophila) (KIRREL3), mRNA
 NM_032536 Homo sapiens netrin G2 (NTNG2), mRNA
 NM_032539 Homo sapiens SLIT and NTRK-like family, member 2 (SLITRK2), mRNA
 NM_032550 Homo sapiens KIAA1914 (KIAA1914), transcript variant 2, mRNA
 NM_032552 Homo sapiens DAB2 interacting protein (DAB2IP), mRNA
 NM_032569 Homo sapiens cytokine-like nuclear factor n-pac (N-PAC), mRNA
 NM_032590 Homo sapiens F-box and leucine-rich repeat protein 10 (FBXL10), mRNA
 NM_032636 Homo sapiens differential display and activated by p53 (DDA3), mRNA
 NM_032869 Homo sapiens chronic myelogenous leukemia tumor antigen 66 (CML66), m
 NM_032870 Homo sapiens chromosome 6 open reading frame 111 (C6orf111), mRNA
 NM_032947 Homo sapiens putative small membrane protein NID67 (NID67), mRNA
 NM_033026 Homo sapiens piccolo (presynaptic cytomatrix protein) (PCLO), transcript va
 NM_033046 Homo sapiens rhotekin (RTKN), mRNA
 NM_033052 Homo sapiens DMRT-like family C2 (DMRTC2), mRNA
 NM_033053 Homo sapiens DMRT-like family C1 (DMRTC1), mRNA
 NM_033055 Homo sapiens hippocampus abundant transcript 1 (HIAT1), mRNA
 NM_033063 Homo sapiens microtubule-associated protein 6 (MAP6), transcript variant 1,
 NM_033064 Homo sapiens ataxia, cerebellar, Cayman type (caytaxin) (ATCAX), mRNA
 NM_033067 Homo sapiens DMRT-like family B with proline-rich C-terminal, 1 (DMRTB1),
 NM_033071 Homo sapiens spectrin repeat containing, nuclear envelope 1 (SYNE1), tra
 NM_033082 Homo sapiens cytokine induced protein 29 kDa (CIP29), mRNA
 NM_033086 Homo sapiens FGD1 family, member 3 (FGD3), mRNA
 NM_033088 Homo sapiens family with sequence similarity 40, member A (FAM40A), mR
 NM_033090 Homo sapiens GREB1 protein (GREB1), transcript variant b, mRNA
 NM_033107 Homo sapiens hypothetical protein BC004923 (LOC85865), mRNA
 NM_033109 Homo sapiens polyribonucleotide nucleotidyltransferase 1 (PNPT1), mRNA
 NM_033112 Homo sapiens chromosome 6 open reading frame 153 (C6orf153), mRNA
 NM_033121 Homo sapiens ankyrin repeat domain 13 (ANKRD13), mRNA
 NM_033129 Homo sapiens scratch homolog 2, zinc finger protein (Drosophila) (SCR2),
 NM_033141 Homo sapiens mitogen-activated protein kinase kinase kinase 9 (MAP3K9), i
 NM_033160 Homo sapiens DKFZP572C163 protein (DKFZP572C163), mRNA
 NM_033161 Homo sapiens surfact 4 (SURF4), mRNA
 NM_033200 Homo sapiens hypothetical protein BC002942 (BC002942), mRNA
 NM_033201 Homo sapiens hypothetical gene BC008967 (BC008967), mRNA
 NM_033206 Homo sapiens hypothetical gene FLJ00060 (FLJ00060), mRNA
 NM_033253 Homo sapiens 5'-nucleotidase, cytosolic IB (NT5C1B), transcript variant 2, m
 NM_033267 Homo sapiens inroquois homeobox protein 2 (IRX2), mRNA

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NM_033271 Homo sapiens BTB (POZ) domain containing 6 (BTBD6), mRNA
 NM_033276 Homo sapiens Ku70-binding protein 3 (KUB3), mRNA
 NM_033288 Homo sapiens zinc finger protein 160 (ZNF160), transcript variant 1, mRNA
 NM_033364 Homo sapiens chromosome 3 open reading frame 15 (C3orf15), mRNA
 NM_033375 Homo sapiens myosin IC (MYO1C), mRNA
 NM_033386 Homo sapiens molecule interacting with Rab13 (MIRAB13), mRNA
 NM_033387 Homo sapiens chromosome 9 open reading frame 59 (C9orf59), mRNA
 NM_033389 Homo sapiens slingshot homolog 2 (Drosophila) (SSH2), mRNA
 NM_033392 Homo sapiens mitogen-activated protein kinase 8 interacting protein 3 (MAPI)
 NM_033393 Homo sapiens KIAA1727 protein (KIAA1727), mRNA
 NM_033396 Homo sapiens tankyrase 1 binding protein 1, 182kDa (TNKS1BP1), mRNA
 NM_033397 Homo sapiens KIAA1754 (KIAA1754), mRNA
 NM_033402 Homo sapiens KIAA1764 protein (KIAA1764), mRNA
 NM_033404 Homo sapiens kinase non-catalytic C-lobe domain (KIND) containing 1 (KND)
 NM_033405 Homo sapiens peroxisomal proliferator-activated receptor A interacting comp
 NM_033407 Homo sapiens dedicator of cytokinesis 7 (DOCK7), mRNA
 NM_033425 Homo sapiens DIX domain containing 1 (DIXDC1), mRNA
 NM_033426 Homo sapiens KIAA1737 (KIAA1737), mRNA
 NM_033429 Homo sapiens calmodulin-like 4 (CALML4), mRNA
 NM_033449 Homo sapiens FCH and double SH3 domains 1 (FCHSD1), mRNA
 NM_033450 Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 10
 NM_033452 Homo sapiens tripartite motif-containing 47 (TRIM47), mRNA
 NM_033505 Homo sapiens selenoprotein I, 1 (SELI), mRNA
 NM_033510 Homo sapiens dispatched homolog 2 (Drosophila) (DISP2), mRNA
 NM_033512 Homo sapiens TSPY-like 5 (TSPYL5), mRNA
 NM_033513 Homo sapiens chromosome 19 open reading frame 20 (C19orf20), mRNA
 NM_033520 Homo sapiens chromosome 19 open reading frame 33 (C19orf33), mRNA
 NM_033542 Homo sapiens chromosome 20 open reading frame 35 (C20orf35), mRNA
 NM_033548 Homo sapiens similar to ZINC FINGER PROTEIN 257 (BONE MARROW ZI
 NM_033553 Homo sapiens guanylate cyclase activator 2A (guanylin) (GUCA2A), mRNA
 NM_033557 Homo sapiens similar to putative transmembrane protein; homolog of yeast C
 NM_033631 Homo sapiens leudne zipper protein 1 (LUZP1), mRNA
 NM_033647 Homo sapiens helicase (DNA) B (HELB), mRNA
 NM_033686 Homo sapiens integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen
 NM_033687 Homo sapiens Integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen
 NM_033688 Homo sapiens Integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen
 NM_033689 Homo sapiens Integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen
 NM_052843 Homo sapiens obscurin, cytoskeletal calmodulin and titin-interacting RhoGEF
 NM_052846 Homo sapiens elastin microfibril interfacer 3 (EMILIN3), mRNA
 NM_052847 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GN
 NM_052849 Homo sapiens hypothetical protein MGC20481 (MGC20481), mRNA
 NM_052850 Homo sapiens growth arrest and DNA-damage-inducible, gamma interacting
 NM_052857 Homo sapiens hypothetical protein MGC20398 (MGC20398), mRNA
 NM_052864 Homo sapiens TRAF2 binding protein (T2BP), mRNA
 NM_052867 Homo sapiens voltage gated channel like 1 (VGCLN1), mRNA
 NM_052878 Homo sapiens chromosome 19 open reading frame 36 (C19orf36), mRNA
 NM_052882 Homo sapiens polycystic kidney disease 1-like 2 (PKD1L2), transcript variant
 NM_052896 Homo sapiens CUB and Sushi multiple domains 2 (CSMD2), mRNA
 NM_052897 Homo sapiens methyl-CpG binding domain protein 6 (MBD6), mRNA
 NM_052899 Homo sapiens G protein-regulated inducer of neurite outgrowth 1 (KIAA1893)
 NM_052900 Homo sapiens CUB and Sushi multiple domains 3 (CSMD3), transcript variant
 NM_052901 Homo sapiens solute carrier family 25 (mitochondrial carrier; phosphate carrier
 NM_052902 Homo sapiens serine/threonine kinase 11 interacting protein (STK11IP), mR
 NM_052903 Homo sapiens tubulin, gamma complex associated protein 5 (TUBGCP5), m
 NM_052904 Homo sapiens KIAA1900 (KIAA1900), mRNA
 NM_052905 Homo sapiens formin-like 2 (FMNL2), mRNA
 NM_052909 Homo sapiens KIAA1909 protein (KIAA1909), mRNA

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NM_130465 Homo sapiens F-box protein 23 (FBXO23), mRNA
 NM_130466 Homo sapiens ubiquitin protein ligase E3B (UBE3B), transcript variant 1, mRNA
 NM_130470 Homo sapiens MAP-kinase activating death domain (MADD), transcript variant 1, mRNA
 NM_130471 Homo sapiens MAP-kinase activating death domain (MADD), transcript variant 2, mRNA
 NM_130472 Homo sapiens MAP-kinase activating death domain (MADD), transcript variant 3, mRNA
 NM_130473 Homo sapiens MAP-kinase activating death domain (MADD), transcript variant 4, mRNA
 NM_130474 Homo sapiens MAP-kinase activating death domain (MADD), transcript variant 5, mRNA
 NM_130475 Homo sapiens MAP-kinase activating death domain (MADD), transcript variant 6, mRNA
 NM_130476 Homo sapiens MAP-kinase activating death domain (MADD), transcript variant 7, mRNA
 NM_130760 Homo sapiens mucosal vascular addressin cell adhesion molecule 1 (MADC), transcript variant 1, mRNA
 NM_130761 Homo sapiens mucosal vascular addressin cell adhesion molecule 1 (MADC), transcript variant 2, mRNA
 NM_130762 Homo sapiens mucosal vascular addressin cell adhesion molecule 1 (MADC), transcript variant 3, mRNA
 NM_130766 Homo sapiens skeletal muscle and kidney enriched inositol phosphatase (SKI), transcript variant 1, mRNA
 NM_130771 Homo sapiens osteoclast-associated receptor (OSCAR), transcript variant 3, mRNA
 NM_130775 Homo sapiens XAGE-5 protein (XAGE-5), mRNA
 NM_130776 Homo sapiens G antigen, family D, 4 (GAGED4), transcript variant 2, mRNA
 NM_130777 Homo sapiens G antigen, family D, 3 (GAGED3), mRNA
 NM_130788 Homo sapiens WW domain containing oxidoreductase (WWOX), transcript variant 1, mRNA
 NM_130790 Homo sapiens WW domain containing oxidoreductase (WWOX), transcript variant 2, mRNA
 NM_130791 Homo sapiens WW domain containing oxidoreductase (WWOX), transcript variant 3, mRNA
 NM_130792 Homo sapiens WW domain containing oxidoreductase (WWOX), transcript variant 4, mRNA
 NM_130793 Homo sapiens nucleolar protein family 6 (RNA-associated) (NOL6), transcript variant 1, mRNA
 NM_130794 Homo sapiens cystatin 11 (CST11), transcript variant 1, mRNA
 NM_130797 Homo sapiens dipeptidylpeptidase 6 (DPP6), transcript variant 1, mRNA
 NM_130798 Homo sapiens synaptosomal-associated protein, 23kDa (SNAP23), transcript variant 1, mRNA
 NM_130799 Homo sapiens multiple endocrine neoplasia I (MEN1), transcript variant 2, mRNA
 NM_130800 Homo sapiens multiple endocrine neoplasia I (MEN1), transcript variant e1B, mRNA
 NM_130801 Homo sapiens multiple endocrine neoplasia I (MEN1), transcript variant e1C, mRNA
 NM_130802 Homo sapiens multiple endocrine neoplasia I (MEN1), transcript variant e1D, mRNA
 NM_130803 Homo sapiens multiple endocrine neoplasia I (MEN1), transcript variant e1E, mRNA
 NM_130804 Homo sapiens multiple endocrine neoplasia I (MEN1), transcript variant e1F, mRNA
 NM_130806 Homo sapiens leucine-rich repeat-containing G protein-coupled receptor 8 (LGR8), transcript variant 1, mRNA
 NM_130807 Homo sapiens MOB1, Mps One Binder kinase activator-like 2A (yeast) (MOB1), transcript variant 1, mRNA
 NM_130808 Homo sapiens copine IV (CPNE4), mRNA
 NM_130809 Homo sapiens hypothetical protein MGC12103 (LOC133619), mRNA
 NM_130810 Homo sapiens dyslexia susceptibility 1 candidate 1 (DYX1C1), mRNA
 NM_130811 Homo sapiens synaptosomal-associated protein, 25kDa (SNAP25), transcript variant 1, mRNA
 NM_130830 Homo sapiens leucine rich repeat containing 15 (LRRC15), mRNA
 NM_130831 Homo sapiens optic atrophy 1 (autosomal dominant) (OPA1), nuclear gene e, mRNA
 NM_130832 Homo sapiens optic atrophy 1 (autosomal dominant) (OPA1), nuclear gene e, mRNA
 NM_130833 Homo sapiens optic atrophy 1 (autosomal dominant) (OPA1), nuclear gene e, mRNA
 NM_130834 Homo sapiens optic atrophy 1 (autosomal dominant) (OPA1), nuclear gene e, mRNA
 NM_130835 Homo sapiens optic atrophy 1 (autosomal dominant) (OPA1), nuclear gene e, mRNA
 NM_130836 Homo sapiens optic atrophy 1 (autosomal dominant) (OPA1), nuclear gene e, mRNA
 NM_130837 Homo sapiens optic atrophy 1 (autosomal dominant) (OPA1), nuclear gene e, mRNA
 NM_130838 Homo sapiens ubiquitin protein ligase E3A (human papilloma virus E6-associ), transcript variant 1, mRNA
 NM_130839 Homo sapiens ubiquitin protein ligase E3A (human papilloma virus E6-associ), transcript variant 2, mRNA
 NM_130840 Homo sapiens ATPase, H⁺ transporting, lysosomal V0 subunit a isoform 4 (ATP5A4), transcript variant 1, mRNA
 NM_130841 Homo sapiens ATPase, H⁺ transporting, lysosomal V0 subunit a isoform 4 (ATP5A4), transcript variant 2, mRNA
 NM_130842 Homo sapiens protein tyrosine phosphatase, receptor type, N polypeptide 2 (PTN2), transcript variant 1, mRNA
 NM_130843 Homo sapiens protein tyrosine phosphatase, receptor type, N polypeptide 2 (PTN2), transcript variant 2, mRNA
 NM_130844 Homo sapiens WW domain containing oxidoreductase (WWOX), transcript variant 1, mRNA
 NM_130845 Homo sapiens syntrophin, beta 2 (dystrophin-associated protein A1, 59kDa), transcript variant 1, mRNA
 NM_130846 Homo sapiens protein tyrosine phosphatase, receptor type, R (PTPRR), transcript variant 1, mRNA
 NM_130847 Homo sapiens angiotensin II type 1 receptor (AGTR1), mRNA
 NM_130848 Homo sapiens dendritic cell nuclear protein 1 (DCNP1), mRNA
 NM_130849 Homo sapiens solute carrier family 39 (zinc transporter), member 4 (SLC39A4), transcript variant 1, mRNA

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NM_130850 Homo sapiens bone morphogenetic protein 4 (BMP4), transcript variant 2, m
 NM_130851 Homo sapiens bone morphogenetic protein 4 (BMP4), transcript variant 3, m
 NM_130852 Homo sapiens palate, lung and nasal epithelium carcinoma associated (PLU
 NM_130853 Homo sapiens protein tyrosine phosphatase, receptor type, S (PTPRS), trans
 NM_130854 Homo sapiens protein tyrosine phosphatase, receptor type, S (PTPRS), trans
 NM_130855 Homo sapiens protein tyrosine phosphatase, receptor type, S (PTPRS), trans
 NM_130856 Homo sapiens WAP four-disulfide core domain 8 (WFDC8), transcript variant
 NM_130857 Homo sapiens dynein, cytoplasmic, light polypeptide 2B (DNCL2B), mRNA
 NM_130858 Homo sapiens cAMP responsive element binding protein 3-like 4 (CREB3L4)
 NM_130899 Homo sapiens hypothetical protein MGC26988 (MGC26988), mRNA
 NM_130900 Homo sapiens retinoic acid early transcript 1L (RAET1L), mRNA
 NM_130901 Homo sapiens chromosome 15 open reading frame 16 (C15orf16), mRNA
 NM_130902 Homo sapiens cytochrome c oxidase subunit VIIb2 (COX7B2), mRNA
 NM_130906 Homo sapiens peptidylprolyl isomerase (cyclophilin)-like 3 (PPII3), transcript
 NM_131915 Homo sapiens similar to hypothetical protein DKFZp434K191 (H. sapiens) (L
 NM_131916 Homo sapiens peptidylprolyl isomerase (cyclophilin)-like 3 (PPII3), transcript
 NM_131917 Homo sapiens Fas (TNFRSF8) associated factor 1 (FAF1), transcript variant
 NM_133168 Homo sapiens osteoclast-associated receptor (OSCAR), transcript variant 5,
 NM_133169 Homo sapiens osteoclast-associated receptor (OSCAR), transcript variant 4,
 NM_133170 Homo sapiens protein tyrosine phosphatase, receptor type, T (PTPRT), trans
 NM_133171 Homo sapiens engulfment and cell motility 2 (ced-12 homolog, C. elegans) (I
 NM_133172 Homo sapiens amyloid beta (A4) precursor protein-binding, family B, membe
 NM_133173 Homo sapiens amyloid beta (A4) precursor protein-binding, family B, membe
 NM_133174 Homo sapiens amyloid beta (A4) precursor protein-binding, family B, membe
 NM_133175 Homo sapiens amyloid beta (A4) precursor protein-binding, family B, membe
 NM_133176 Homo sapiens amyloid beta (A4) precursor protein-binding, family B, membe
 NM_133177 Homo sapiens protein tyrosine phosphatase, receptor type, U (PTPRU), trans
 NM_133178 Homo sapiens protein tyrosine phosphatase, receptor type, U (PTPRU), trans
 NM_133179 Homo sapiens G antigen, family D, 4 (GAGED4), transcript variant 1, mRNA
 NM_133180 Homo sapiens EPS8-like 1 (EPS8L1), transcript variant 1, mRNA
 NM_133181 Homo sapiens EPS8-like 3 (EPS8L3), transcript variant 2, mRNA
 NM_133259 Homo sapiens leucine-rich PPR-motif containing (LRPPRC), mRNA
 NM_133261 Homo sapiens PDZ domain protein GIPC3 (GIPC3), mRNA
 NM_133262 Homo sapiens ATPase, H⁺ transporting, lysosomal 13kDa, V1 subunit G Isol
 NM_133263 Homo sapiens peroxisome proliferative activated receptor, gamma, coactivat
 NM_133264 Homo sapiens WIRE protein (WIRE), mRNA
 NM_133265 Homo sapiens angiotensin (AMOT), mRNA
 NM_133266 Homo sapiens SH3 and multiple ankyrin repeat domains 2 (SHANK2), trans
 NM_133267 Homo sapiens homeobox protein GSH-2 (GSH-2), mRNA
 NM_133268 Homo sapiens oxysterol binding protein-like 1A (OSBPL1A), transcript varian
 NM_133269 Homo sapiens Fc fragment of IgA, receptor for (FCAR), transcript variant 2, r
 NM_133271 Homo sapiens Fc fragment of IgA, receptor for (FCAR), transcript variant 3, r
 NM_133272 Homo sapiens Fc fragment of IgA, receptor for (FCAR), transcript variant 4, r
 NM_133273 Homo sapiens Fc fragment of IgA, receptor for (FCAR), transcript variant 5, r
 NM_133274 Homo sapiens Fc fragment of IgA, receptor for (FCAR), transcript variant 6, r
 NM_133277 Homo sapiens Fc fragment of IgA, receptor for (FCAR), transcript variant 7, r
 NM_133278 Homo sapiens Fc fragment of IgA, receptor for (FCAR), transcript variant 8, r
 NM_133279 Homo sapiens Fc fragment of IgA, receptor for (FCAR), transcript variant 9, r
 NM_133280 Homo sapiens Fc fragment of IgA, receptor for (FCAR), transcript variant 10,
 NM_133282 Homo sapiens RAD1 homolog (S. pombe) (RAD1), transcript variant 2, mRN
 NM_133325 Homo sapiens PHD finger protein 10 (PHF10), transcript variant 2, mRNA
 NM_133326 Homo sapiens ATPase, H⁺ transporting, lysosomal 13kDa, V1 subunit G Isol
 NM_133327 Homo sapiens sema domain, transmembrane domain (TM), and cytoplasmic
 NM_133328 Homo sapiens death effector domain containing 2 (DEDD2), mRNA
 NM_133329 Homo sapiens potassium voltage-gated channel, subfamily G, member 3 (KC
 NM_133330 Homo sapiens Wolf-Hirschhorn syndrome candidate 1 (WHSC1), transcript v
 NM_133331 Homo sapiens Wolf-Hirschhorn syndrome candidate 1 (WHSC1), transcript v

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NM_133332 Homo sapiens Wolf-Hirschhorn syndrome candidate 1 (WHSC1), transcript \
 NM_133333 Homo sapiens Wolf-Hirschhorn syndrome candidate 1 (WHSC1), transcript \
 NM_133334 Homo sapiens Wolf-Hirschhorn syndrome candidate 1 (WHSC1), transcript \
 NM_133335 Homo sapiens Wolf-Hirschhorn syndrome candidate 1 (WHSC1), transcript \
 NM_133336 Homo sapiens Wolf-Hirschhorn syndrome candidate 1 (WHSC1), transcript \
 NM_133337 Homo sapiens fer-1-like 3, myoferlin (C. elegans) (FER1L3), transcript varian
 NM_133338 Homo sapiens RAD17 homolog (S. pombe) (RAD17), transcript variant 1, mf
 NM_133339 Homo sapiens RAD17 homolog (S. pombe) (RAD17), transcript variant 2, mf
 NM_133340 Homo sapiens RAD17 homolog (S. pombe) (RAD17), transcript variant 3, mf
 NM_133341 Homo sapiens RAD17 homolog (S. pombe) (RAD17), transcript variant 4, mf
 NM_133342 Homo sapiens RAD17 homolog (S. pombe) (RAD17), transcript variant 5, mf
 NM_133343 Homo sapiens RAD17 homolog (S. pombe) (RAD17), transcript variant 6, mf
 NM_133344 Homo sapiens RAD17 homolog (S. pombe) (RAD17), transcript variant 7, mf
 NM_133367 Homo sapiens chromosome 8 open reading frame 33 (C6orf33), mRNA
 NM_133368 Homo sapiens KIAA1972 protein (KIAA1972), mRNA
 NM_133370 Homo sapiens splicing factor YT521-B (YT521), mRNA
 NM_133371 Homo sapiens myozenin 3 (MYOZ3), mRNA
 NM_133373 Homo sapiens phospholipase C, delta 3 (PLCD3), mRNA
 NM_133375 Homo sapiens hypothetical protein MGC4562 (MGC4562), mRNA
 NM_133376 Homo sapiens integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen
 NM_133377 Homo sapiens RAD1 homolog (S. pombe) (RAD1), transcript variant 3, mRNA
 NM_133378 Homo sapiens titin (TTN), transcript variant N2-A, mRNA
 NM_133379 Homo sapiens titin (TTN), transcript variant novex-3, mRNA
 NM_133430 Homo sapiens G antigen, family D, 2 (GAGED2), transcript variant 3, mRNA
 NM_133431 Homo sapiens G antigen, family D, 2 (GAGED2), transcript variant 2, mRNA
 NM_133432 Homo sapiens titin (TTN), transcript variant novex-1, mRNA
 NM_133433 Homo sapiens Nipped-B homolog (Drosophila) (NIPBL), transcript variant A,
 NM_133436 Homo sapiens asparagine synthetase (ASNS), transcript variant 1, mRNA
 NM_133437 Homo sapiens titin (TTN), transcript variant novex-2, mRNA
 NM_133439 Homo sapiens transcriptional adaptor 2 (ADA2 homolog, yeast)-like (TADA2)
 NM_133443 Homo sapiens glutamic pyruvate transaminase (alanine aminotransferase) 2
 NM_133444 Homo sapiens zinc finger protein 526 (ZNF526), mRNA
 NM_133445 Homo sapiens glutamate receptor, ionotropic, N-methyl-D-aspartate 3A (GRI
 NM_133446 Homo sapiens centaurin, gamma-like family, member 1 (CTGLF1), mRNA
 NM_133448 Homo sapiens KIAA1944 protein (KIAA1944), mRNA
 NM_133450 Homo sapiens KIAA1977 protein (KIAA1977), mRNA
 NM_133452 Homo sapiens RAVR1 (RAVER1), mRNA
 NM_133455 Homo sapiens EMI domain containing 1 (EMID1), mRNA
 NM_133456 Homo sapiens apical protein 2 (APXL2), mRNA
 NM_133457 Homo sapiens EMI domain containing 2 (EMID2), mRNA
 NM_133459 Homo sapiens KIAA1983 protein (FLJ30881), mRNA
 NM_133462 Homo sapiens tetratricopeptide repeat domain 14 (TTC14), mRNA
 NM_133466 Homo sapiens zinc finger protein 545 (ZNF545), mRNA
 NM_133467 Homo sapiens Cbp/p300-interacting transactivator, with Glu/Asp-rich carbox
 NM_133468 Homo sapiens BMP-binding endothelial regulator precursor protein (BMPER),
 NM_133473 Homo sapiens zinc finger protein 431 (ZNF431), mRNA
 NM_133474 Homo sapiens KIAA1982 protein (KIAA1982), mRNA
 NM_133476 Homo sapiens zinc finger protein 384 (ZNF384), mRNA
 NM_133478 Homo sapiens solute carrier family 4, sodium bicarbonate cotransporter, mer
 NM_133479 Homo sapiens solute carrier family 4, sodium bicarbonate cotransporter, mer
 NM_133480 Homo sapiens transcriptional adaptor 3 (NGG1 homolog, yeast)-like (TADA3)
 NM_133481 Homo sapiens transcriptional adaptor 3 (NGG1 homolog, yeast)-like (TADA3)
 NM_133482 Homo sapiens RAD50 homolog (S. cerevisiae) (RAD50), transcript variant 2,
 NM_133483 Homo sapiens RAC/CDC42 exchange factor (GEFT), transcript variant 2, mf
 NM_133484 Homo sapiens TRAF family member-associated NFkB activator (TANK), trar
 NM_133486 Homo sapiens muscleblind-like 3 (Drosophila) (MBNL3), mRNA
 NM_133487 Homo sapiens RAD51 homolog (RecA homolog, E. coli) (S. cerevisiae) (RAC

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NM_133489 Homo sapiens solute carrier family 26, member 10 (SLC26A10), mRNA
 NM_133490 Homo sapiens potassium voltage-gated channel, subfamily G, member 4 (KC
 NM_133491 Homo sapiens spermidine/spermine N1-acetyltransferase 2 (SAT2), mRNA
 NM_133492 Homo sapiens N-acylsphingosine amidohydrolase (alkaline ceramidase) 3 (A
 NM_133493 Homo sapiens CD109 antigen (Gov platelet alloantigens) (CD109), mRNA
 NM_133494 Homo sapiens NIMA (never in mitosis gene a)-related kinase 7 (NEK7), mRNA
 NM_133496 Homo sapiens solute carrier family 30 (zinc transporter), member 7 (SLC30A
 NM_133497 Homo sapiens potassium channel, subfamily V, member 2 (KCNV2), mRNA
 NM_133498 Homo sapiens sperm acrosome associated 4 (SPAC4), mRNA
 NM_133499 Homo sapiens synapsin I (SYN1), transcript variant 1b, mRNA
 NM_133502 Homo sapiens zinc finger protein 274 (ZNF274), transcript variant ZNF274c,
 NM_133503 Homo sapiens decorin (DCN), transcript variant A2, mRNA
 NM_133504 Homo sapiens decorin (DCN), transcript variant B, mRNA
 NM_133505 Homo sapiens decorin (DCN), transcript variant C, mRNA
 NM_133506 Homo sapiens decorin (DCN), transcript variant D, mRNA
 NM_133507 Homo sapiens decorin (DCN), transcript variant E, mRNA
 NM_133509 Homo sapiens RAD51-like 1 (S. cerevisiae) (RAD51L1), transcript variant 3,
 NM_133510 Homo sapiens RAD51-like 1 (S. cerevisiae) (RAD51L1), transcript variant 2,
 NM_133625 Homo sapiens synapsin II (SYN2), transcript variant 1la, mRNA
 NM_133627 Homo sapiens RAD51-like 3 (S. cerevisiae) (RAD51L3), transcript variant 2,
 NM_133628 Homo sapiens RAD51-like 3 (S. cerevisiae) (RAD51L3), transcript variant 3,
 NM_133629 Homo sapiens RAD51-like 3 (S. cerevisiae) (RAD51L3), transcript variant 4,
 NM_133630 Homo sapiens RAD51-like 3 (S. cerevisiae) (RAD51L3), transcript variant 5,
 NM_133631 Homo sapiens roundabout, axon guidance receptor, homolog 1 (Drosophila)
 NM_133632 Homo sapiens synapsin III (SYN3), transcript variant 1Iib, mRNA
 NM_133633 Homo sapiens synapsin III (SYN3), transcript variant 1Iic, mRNA
 NM_133634 Homo sapiens protein O-fucosyltransferase 2 (POFUT2), transcript variant 2,
 NM_133635 Homo sapiens protein O-fucosyltransferase 2 (POFUT2), transcript variant 3,
 NM_133636 Homo sapiens DNA helicase HEL308 (HEL308), mRNA
 NM_133637 Homo sapiens DEAQ box polypeptide 1 (RNA-dependent ATPase) (DQX1), I
 NM_133638 Homo sapiens a disintegrin-like and metalloprotease (repolyisin type) with th
 NM_133639 Homo sapiens ras homolog gene family, member V (RHOF), mRNA
 NM_133640 Homo sapiens surfeit 5 (SURF5), transcript variant b, mRNA
 NM_133642 Homo sapiens like-glycosyltransferase (LARGE), transcript variant 2, mRNA
 NM_133644 Homo sapiens GTP binding protein 3 (mitochondrial) (GTPBP3), mRNA
 NM_133645 Homo sapiens mitochondrial translation optimization 1 homolog (S. cerevisia
 NM_133646 Homo sapiens sterile alpha motif and leucine zipper containing kinase AZK (A
 NM_133650 Homo sapiens spectrin repeat containing, nuclear envelope 1 (SYNE1), trans
 NM_134258 Homo sapiens transducin (beta)-like 1Y-linked (TBL1Y), transcript variant 2, I
 NM_134259 Homo sapiens transducin (beta)-like 1Y-linked (TBL1Y), transcript variant 3, I
 NM_134260 Homo sapiens RAR-related orphan receptor A (RORA), transcript variant 2, r
 NM_134261 Homo sapiens RAR-related orphan receptor A (RORA), transcript variant 1, r
 NM_134262 Homo sapiens RAR-related orphan receptor A (RORA), transcript variant 4, r
 NM_134263 Homo sapiens solute carrier family 26, member 6 (SLC26A6), transcript vari
 NM_134264 Homo sapiens WD repeat and SOCS box-containing 1 (WSB1), transcript va
 NM_134265 Homo sapiens WD repeat and SOCS box-containing 1 (WSB1), transcript va
 NM_134266 Homo sapiens solute carrier family 26, member 7 (SLC26A7), transcript vari
 NM_134268 Homo sapiens cytoglobin (CYGB), mRNA
 NM_134269 Homo sapiens smoothenin (SMTN), transcript variant 2, mRNA
 NM_134270 Homo sapiens smoothenin (SMTN), transcript variant 1, mRNA
 NM_134323 Homo sapiens TAR (HIV) RNA binding protein 2 (TARBP2), transcript varian
 NM_134324 Homo sapiens TAR (HIV) RNA binding protein 2 (TARBP2), transcript varian
 NM_134325 Homo sapiens solute carrier family 26, member 9 (SLC26A9), transcript vari
 NM_134421 Homo sapiens hippocampal-like 1 (HPCAL1), transcript variant 2, mRNA
 NM_134422 Homo sapiens RAD52 homolog (S. cerevisiae) (RAD52), transcript variant de
 NM_134423 Homo sapiens RAD52 homolog (S. cerevisiae) (RAD52), transcript variant ge
 NM_134424 Homo sapiens RAD52 homolog (S. cerevisiae) (RAD52), transcript variant be

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NM_134425 Homo sapiens solute carrier family 26 (sulfate transporter), member 1 (SLC2
 NM_134426 Homo sapiens solute carrier family 26, member 6 (SLC26A6), transcript vari
 NM_134427 Homo sapiens regulator of G-protein signalling 3 (RGS3), transcript variant 4
 NM_134428 Homo sapiens regulatory factor X, 3 (influences HLA class II expression) (RF
 NM_134431 Homo sapiens solute carrier organic anion transporter family, member 1A2 (S
 NM_134433 Homo sapiens regulatory factor X, 2 (influences HLA class II expression) (RF
 NM_134434 Homo sapiens RAD54 homolog B (S. cerevisiae) (RAD54B), transcript varian
 NM_134440 Homo sapiens regulatory factor X-associated ankyrin-containing protein (RF)
 NM_134441 Homo sapiens relaxin 2 (h2) (RLN2), transcript variant 1, mRNA
 NM_134442 Homo sapiens cAMP responsive element binding protein 1 (CREB1), transcr
 NM_134444 Homo sapiens NACHT, leucine rich repeat and PYD containing 4 (NALP4), n
 NM_134445 Homo sapiens CD99 antigen-like 2 (CD99L2), mRNA
 NM_134446 Homo sapiens CD99 antigen-like 2 (CD99L2), mRNA
 NM_134447 Homo sapiens chromosome 19 open reading frame 2 (C19orf2), transcript vs
 NM_134470 Homo sapiens Interleukin 1 receptor accessory protein (IL1RAP), transcript v
 NM_138270 Homo sapiens alpha thalassemia/mental retardation syndrome X-linked (RAI
 NM_138271 Homo sapiens alpha thalassemia/mental retardation syndrome X-linked (RAI
 NM_138272 Homo sapiens chromosome 6 open reading frame 25 (C6orf25), transcript vs
 NM_138273 Homo sapiens chromosome 6 open reading frame 25 (C6orf25), transcript vs
 NM_138274 Homo sapiens chromosome 6 open reading frame 25 (C6orf25), transcript vs
 NM_138275 Homo sapiens chromosome 6 open reading frame 25 (C6orf25), transcript vs
 NM_138276 Homo sapiens chromosome 6 open reading frame 25 (C6orf25), transcript vs
 NM_138277 Homo sapiens chromosome 6 open reading frame 25 (C6orf25), transcript vs
 NM_138278 Homo sapiens BCL2/adenovirus E1B 19kD interacting protein like (BNIPL), r
 NM_138279 Homo sapiens BCL2/adenovirus E1B 19kD interacting protein like (BNIPL), r
 NM_138280 Homo sapiens citrate lyase beta like (CLYBL), transcript variant 1, mRNA
 NM_138281 Homo sapiens distal-less homeobox 4 (DLX4), transcript variant 1, mRNA
 NM_138282 Homo sapiens ATPase, H+ transporting, lysosomal 13kDa, V1 subunit G iso
 NM_138283 Homo sapiens cystatin-like 1 (CSTL1), mRNA
 NM_138284 Homo sapiens Interleukin 17D (IL17D), mRNA
 NM_138285 Homo sapiens nucleoporin 35kDa (NUP35), mRNA
 NM_138286 Homo sapiens hypothetical protein FLJ31526 (LOC148213), mRNA
 NM_138287 Homo sapiens rhysin 2 (BBAP), mRNA
 NM_138288 Homo sapiens chromosome 14 open reading frame 147 (C14orf147), mRNA
 NM_138289 Homo sapiens actin-related protein T1 (ACTRT1), mRNA
 NM_138290 Homo sapiens Rap2-binding protein 9 (RPIB9), mRNA
 NM_138292 Homo sapiens ataxia telangiectasia mutated (includes complementation gro
 NM_138293 Homo sapiens ataxia telangiectasia mutated (includes complementation gro
 NM_138294 Homo sapiens expressed in prostate and testis (PATE), mRNA
 NM_138295 Homo sapiens polycystic kidney disease 1 like 1 (PKD1L1), mRNA
 NM_138296 Homo sapiens pre T-cell antigen receptor alpha (PTCRA), mRNA
 NM_138297 Homo sapiens mucin 4, tracheobronchial (MUC4), transcript variant 5, mRNA
 NM_138298 Homo sapiens mucin 4, tracheobronchial (MUC4), transcript variant 2, mRNA
 NM_138299 Homo sapiens mucin 4, tracheobronchial (MUC4), transcript variant 3, mRNA
 NM_138300 Homo sapiens pygopus 2 (PYGO2), mRNA
 NM_138316 Homo sapiens pantothenate kinase 1 (PANK1), transcript variant gamma, m
 NM_138317 Homo sapiens potassium channel, subfamily K, member 10 (KCNK10), trans
 NM_138318 Homo sapiens potassium channel, subfamily K, member 10 (KCNK10), trans
 NM_138319 Homo sapiens proprotein convertase subtilisin/kexin type 6 (PCSK6), transcr
 NM_138320 Homo sapiens proprotein convertase subtilisin/kexin type 6 (PCSK6), transcr
 NM_138321 Homo sapiens proprotein convertase subtilisin/kexin type 6 (PCSK6), transcr
 NM_138322 Homo sapiens proprotein convertase subtilisin/kexin type 6 (PCSK6), transcr
 NM_138323 Homo sapiens proprotein convertase subtilisin/kexin type 6 (PCSK6), transcr
 NM_138324 Homo sapiens proprotein convertase subtilisin/kexin type 6 (PCSK6), transcr
 NM_138325 Homo sapiens proprotein convertase subtilisin/kexin type 6 (PCSK6), transcr
 NM_138326 Homo sapiens aminocarboxymuconate semialdehyde decarboxylase (ACMS
 NM_138327 Homo sapiens trace amine receptor 1 (TRAR1), mRNA

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NM_138328 Homo sapiens rhomboid, veinlet-like 4 (Drosophila) (RHBDL4), mRNA
 NM_138329 Homo sapiens NACHT, leucine rich repeat and PYD containing 6 (NALP6), n
 NM_138330 Homo sapiens TRAF6-inhibitory zinc finger protein (TI2), mRNA
 NM_138331 Homo sapiens ribonuclease, RNase A family, 8 (RNASE8), mRNA
 NM_138333 Homo sapiens chromosome 9 open reading frame 42 (C9orf42), mRNA
 NM_138344 Homo sapiens hypothetical transmembrane protein SBB154 (SBB154), mRNA
 NM_138335 Homo sapiens glucosamine-6-phosphate deaminase 2 (GNPDA2), mRNA
 NM_138336 Homo sapiens helicase/primease complex protein (LOC150678), mRNA
 NM_138337 Homo sapiens myeloid inhibitory C-type lectin-like receptor (MIL), transcript
 NM_138338 Homo sapiens polymerase (RNA) III (DNA directed) polypeptide H (22.9kD) (n
 NM_138340 Homo sapiens abhydrolase domain containing 3 (ABHD3), mRNA
 NM_138341 Homo sapiens hypothetical protein BC000282 (LOC89894), mRNA
 NM_138342 Homo sapiens hypothetical protein BC008326 (LOC89944), mRNA
 NM_138343 Homo sapiens kinesin-like 8 (KNSL8), transcript variant 4, mRNA
 NM_138344 Homo sapiens chromosome 14 open reading frame 152 (C14orf152), mRNA
 NM_138346 Homo sapiens hypothetical protein MGC33867 (MGC33867), mRNA
 NM_138347 Homo sapiens zinc finger protein 551 (ZNF551), mRNA
 NM_138348 Homo sapiens hypothetical protein BC007706 (LOC90268), mRNA
 NM_138349 Homo sapiens hypothetical protein BC004507 (LOC90313), mRNA
 NM_138350 Homo sapiens hypothetical protein MGC33488 (MGC33488), mRNA
 NM_138352 Homo sapiens atherin (LOC90378), mRNA
 NM_138355 Homo sapiens secernin 2 (Ses2), mRNA
 NM_138356 Homo sapiens hypothetical protein BC007586 (LOC90525), mRNA
 NM_138357 Homo sapiens chromosome 10 open reading frame 42 (C10orf42), mRNA
 NM_138358 Homo sapiens hypothetical protein BC011833 (LOC90580), mRNA
 NM_138360 Homo sapiens hypothetical protein BC008134 (LOC90668), mRNA
 NM_138361 Homo sapiens leucine rich repeat and sterile alpha motif containing 1 (LRSAn
 NM_138362 Homo sapiens hypothetical protein BC000919 (LOC90736), mRNA
 NM_138363 Homo sapiens hypothetical protein BC009518 (LOC90799), mRNA
 NM_138364 Homo sapiens hypothetical protein BC004337 (LOC90826), mRNA
 NM_138368 Homo sapiens hypothetical protein BC004895 (LOC91056), mRNA
 NM_138369 Homo sapiens family with sequence similarity 44, member B (FAM44B), mRn
 NM_138371 Homo sapiens hypothetical protein MGC16044 (MGC16044), mRNA
 NM_138372 Homo sapiens hypothetical protein BC001610 (LOC91661), mRNA
 NM_138373 Homo sapiens myeloid-associated differentiation marker (MYADM), mRNA
 NM_138375 Homo sapiens Cdk5 and Abl enzyme substrate 1 (CABLES1), mRNA
 NM_138376 Homo sapiens tetratricopeptide repeat domain 5 (TTCS), mRNA
 NM_138379 Homo sapiens T-cell Immunoglobulin and mucin domain containing 4 (TIMD4
 NM_138381 Homo sapiens hypothetical protein BC008322 (MGC15763), mRNA
 NM_138383 Homo sapiens hypothetical protein BC002770 (LOC92154), mRNA
 NM_138384 Homo sapiens shadow of prion protein (Sprn), mRNA
 NM_138385 Homo sapiens hypothetical protein BC009331 (LOC92305), mRNA
 NM_138386 Homo sapiens hypothetical protein BC008207 (LOC92345), mRNA
 NM_138387 Homo sapiens glucose 6 phosphatase, catalytic, 3 (G6PC3), mRNA
 NM_138389 Homo sapiens hypothetical protein BC001096 (LOC92689), mRNA
 NM_138390 Homo sapiens hypothetical protein BC008604 (LOC92691), mRNA
 NM_138391 Homo sapiens chromosome 1 open reading frame 37 (C1orf37), mRNA
 NM_138392 Homo sapiens hypothetical protein BC007853 (LOC92799), mRNA
 NM_138393 Homo sapiens chromosome 19 open reading frame 32 (C19orf32), mRNA
 NM_138394 Homo sapiens hypothetical protein BC008217 (LOC92906), mRNA
 NM_138395 Homo sapiens mitochondrial methionyl-tRNA synthetase (MetRS), mRNA
 NM_138396 Homo sapiens hypothetical protein BC009489 (LOC92979), mRNA
 NM_138397 Homo sapiens hypothetical protein BC012317 (LOC93082), mRNA
 NM_138399 Homo sapiens hypothetical protein BC007772 (LOC93109), mRNA
 NM_138401 Homo sapiens hypothetical protein BC011840 (LOC93343), mRNA
 NM_138402 Homo sapiens hypothetical protein BC004921 (LOC93349), mRNA
 NM_138403 Homo sapiens myosin light chain 2, precursor lymphocyte-specific (MYLC2P

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NM_138408 Homo sapiens chromosome 6 open reading frame 51 (C6orf51), mRNA
 NM_138409 Homo sapiens chromosome 6 open reading frame 117 (C6orf117), mRNA
 NM_138410 Homo sapiens chemokine-like factor super family 7 (CKLFSF7), transcript va
 NM_138412 Homo sapiens retinol dehydrogenase 13 (all-trans and 9-cis) (RDH13), mRN
 NM_138413 Homo sapiens chromosome 10 open reading frame 65 (C10orf65), mRNA
 NM_138414 Homo sapiens hypothetical protein BC011981 (LOC112869), mRNA
 NM_138415 Homo sapiens hypothetical protein BC012187 (LOC112885), mRNA
 NM_138416 Homo sapiens hypothetical protein BC011001 (LOC112937), mRNA
 NM_138417 Homo sapiens hypothetical protein BC012173 (MGC20419), mRNA
 NM_138418 Homo sapiens hypothetical protein MGC15416 (MGC15416), mRNA
 NM_138419 Homo sapiens DUF729 domain containing 1 (DUF71), mRNA
 NM_138421 Homo sapiens hypothetical protein BC012010 (LOC113174), mRNA
 NM_138422 Homo sapiens hypothetical protein BC011824 (LOC113179), mRNA
 NM_138423 Homo sapiens H63 breast cancer expressed gene (H63), transcript variant 1
 NM_138424 Homo sapiens kinesin family member 12 (KIF12), mRNA
 NM_138425 Homo sapiens likely ortholog of mouse gene rich cluster, C10 gene (GRCC1
 NM_138428 Homo sapiens hypothetical protein BC011880 (LOC113444), mRNA
 NM_138429 Homo sapiens claudin 15 (CLDN15), mRNA
 NM_138430 Homo sapiens ADP-ribosylhydrolase like 1 (ADPRHL1), transcript variant 1,
 NM_138431 Homo sapiens hypothetical protein BC011982 (LOC113655), mRNA
 NM_138432 Homo sapiens serine dehydratase-like (SDSL), mRNA
 NM_138433 Homo sapiens hypothetical protein BC009980 (MGC16635), mRNA
 NM_138434 Homo sapiens chromosome 7 open reading frame 29 (C7orf29), mRNA
 NM_138435 Homo sapiens hypothetical protein BC011204 (LOC113828), mRNA
 NM_138436 Homo sapiens hypothetical protein BC013035 (LOC114926), mRNA
 NM_138437 Homo sapiens GASP2 protein (GASP2), mRNA
 NM_138439 Homo sapiens hypothetical protein BC014089 (LOC114984), mRNA
 NM_138440 Homo sapiens hypothetical protein BC013767 (LOC114990), mRNA
 NM_138441 Homo sapiens chromosome 6 open reading frame 150 (C6orf150), mRNA
 NM_138442 Homo sapiens hypothetical protein BC013949 (LOC115098), mRNA
 NM_138443 Homo sapiens coiled-coil domain containing 5 (spindle associated) (CCDC5)
 NM_138444 Homo sapiens potassium channel tetramerisation domain containing 12 (KC
 NM_138445 Homo sapiens G protein-coupled receptor 146 (GPR146), mRNA
 NM_138446 Homo sapiens chromosome 7 open reading frame 30 (C7orf30), mRNA
 NM_138447 Homo sapiens hypothetical protein BC014000 (LOC115509), mRNA
 NM_138448 Homo sapiens acylphosphatase 2, muscle type (ACYP2), mRNA
 NM_138450 Homo sapiens ADP-ribosylation factor-like 11 (ARL11), mRNA
 NM_138451 Homo sapiens hypothetical protein BC013151 (LOC115811), mRNA
 NM_138452 Homo sapiens dehydrogenase/reductase (SDR family) member 1 (DHR51),
 NM_138453 Homo sapiens RAB3C, member RAS oncogene family (RAB3C), mRNA
 NM_138454 Homo sapiens thioredoxin-like 6 (TXNL6), mRNA
 NM_138455 Homo sapiens collagen triple helix repeat containing 1 (CTHRC1), mRNA
 NM_138456 Homo sapiens hypothetical protein BC012330 (MGC20410), mRNA
 NM_138457 Homo sapiens forkhead box P4 (FOXP4), mRNA
 NM_138458 Homo sapiens hypothetical protein BC014022 (LOC116143), mRNA
 NM_138459 Homo sapiens chromosome 6 open reading frame 68 (C6orf68), mRNA
 NM_138460 Homo sapiens chemokine-like factor super family 5 (CKLFSF5), transcript va
 NM_138461 Homo sapiens hypothetical protein BC013113 (LOC116211), mRNA
 NM_138462 Homo sapiens zinc finger, MYND domain containing 19 (ZMYND19), mRNA
 NM_138463 Homo sapiens hypothetical protein BC014072 (LOC116238), mRNA
 NM_138465 Homo sapiens GLI-Kruppel family member GLI4 (GLI4), mRNA
 NM_138467 Homo sapiens hypothetical protein BC009514 (LOC127253), mRNA
 NM_138468 Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region,
 NM_138471 Homo sapiens hypothetical protein BC007540 (LOC144097), mRNA
 NM_138473 Homo sapiens Sp1 transcription factor (SP1), mRNA
 NM_138476 Homo sapiens hypothetical protein MGC5987 (MGC5987), mRNA
 NM_138477 Homo sapiens congenital dyserythropoietic anemia, type I (CDAN1), mRNA

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NM_138479 Homo sapiens hypothetical protein BC007899 (LOC148898), mRNA
 NM_138482 Homo sapiens hypothetical protein BC009264 (LOC151534), mRNA
 NM_138484 Homo sapiens shugoshin-like 1 (S. pombe) (SGOL1), mRNA
 NM_138487 Homo sapiens hypothetical protein BC007882 (LOC152217), mRNA
 NM_138492 Homo sapiens hypothetical protein MGC21644 (MGC21644), transcript varia
 NM_138494 Homo sapiens vav-1 interacting Kruppel-like protein (VIK), transcript variant
 NM_138497 Homo sapiens hypothetical protein BC008050 (LOC158435), mRNA
 NM_138499 Homo sapiens PWWP domain containing 2 (PWWP2), mRNA
 NM_138501 Homo sapiens glycoprotein, synaptic 2 (GPSN2), mRNA
 NM_138551 Homo sapiens thymic stromal lymphopoietin (TSLP), transcript variant 2, mR
 NM_138553 Homo sapiens B-cell CLL/lymphoma 11A (zinc finger protein) (BCL11A), tran
 NM_138554 Homo sapiens toll-like receptor 4 (TLR4), transcript variant 1, mRNA
 NM_138555 Homo sapiens kinesin family member 23 (KIF23), transcript variant 1, mRNA
 NM_138556 Homo sapiens toll-like receptor 4 (TLR4), transcript variant 2, mRNA
 NM_138557 Homo sapiens toll-like receptor 4 (TLR4), transcript variant 4, mRNA
 NM_138558 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 8 (PPP1F
 NM_138559 Homo sapiens B-cell CLL/lymphoma 11A (zinc finger protein) (BCL11A), tran
 NM_138563 Homo sapiens kallikrein 15 (KLK15), transcript variant 2, mRNA
 NM_138564 Homo sapiens kallikrein 15 (KLK15), transcript variant 3, mRNA
 NM_138565 Homo sapiens cortactin (CTTN), transcript variant 2, mRNA
 NM_138566 Homo sapiens glutaminase 2 (liver, mitochondrial) (GLS2), nuclear gene enc
 NM_138567 Homo sapiens synaptotagmin VIII (SYT8), mRNA
 NM_138568 Homo sapiens protein 7 transactivated by hepatitis B virus X antigen (HBxAg
 NM_138569 Homo sapiens chromosome 6 open reading frame 142 (C6orf142), mRNA
 NM_138570 Homo sapiens hypothetical protein MGC15523 (MGC15523), mRNA
 NM_138571 Homo sapiens histidine triad nucleotide binding protein 3 (HINT3), mRNA
 NM_138572 Homo sapiens taube nuss homolog (mouse) (TBN), mRNA
 NM_138573 Homo sapiens neuregulin 4 (LOC145957), mRNA
 NM_138574 Homo sapiens PWWP domain containing 1 (PWWP1), mRNA
 NM_138575 Homo sapiens hypothetical protein MGC5352 (MGC5352), mRNA
 NM_138576 Homo sapiens B-cell CLL/lymphoma 11B (zinc finger protein) (BCL11B), tran
 NM_138578 Homo sapiens BCL2-like 1 (BCL2L1), nuclear gene encoding mitochondrial
 NM_138608 Homo sapiens metallophosphoesterase 1 (MPPE1), mRNA
 NM_138609 Homo sapiens H2A histone family, member Y (H2AFY), transcript variant 1, i
 NM_138610 Homo sapiens H2A histone family, member Y (H2AFY), transcript variant 3, i
 NM_138612 Homo sapiens hyaluronan synthase 3 (HAS3), transcript variant 2, mRNA
 NM_138614 Homo sapiens DEAH (Asp-Glu-Ala-His) box polypeptide 30 (DHX30), transci
 NM_138615 Homo sapiens DEAH (Asp-Glu-Ala-His) box polypeptide 30 (DHX30), transci
 NM_138616 Homo sapiens Rhesus blood group, CcEe antigens (RHCE), transcript varia
 NM_138617 Homo sapiens Rhesus blood group, CcEe antigens (RHCE), transcript varia
 NM_138618 Homo sapiens Rhesus blood group, CcEe antigens (RHCE), transcript varia
 NM_138619 Homo sapiens golgi associated, gamma adaptin ear containing, ARF binding
 NM_138620 Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 31 (DDX31), transc
 NM_138621 Homo sapiens BCL2-like 11 (apoptosis facilitator) (BCL2L11), transcript vari
 NM_138622 Homo sapiens BCL2-like 11 (apoptosis facilitator) (BCL2L11), transcript vari
 NM_138623 Homo sapiens BCL2-like 11 (apoptosis facilitator) (BCL2L11), transcript vari
 NM_138624 Homo sapiens BCL2-like 11 (apoptosis facilitator) (BCL2L11), transcript vari
 NM_138625 Homo sapiens BCL2-like 11 (apoptosis facilitator) (BCL2L11), transcript vari
 NM_138626 Homo sapiens BCL2-like 11 (apoptosis facilitator) (BCL2L11), transcript vari
 NM_138627 Homo sapiens BCL2-like 11 (apoptosis facilitator) (BCL2L11), transcript vari
 NM_138632 Homo sapiens Tara-like protein (HRIHFB2122), transcript variant 2, mRNA
 NM_138633 Homo sapiens A kinase (PRKA) anchor protein 7 (AKAP7), transcript variant
 NM_138634 Homo sapiens microseminoprotein, beta- (MSMB), transcript variant PSP57,
 NM_138635 Homo sapiens H2A histone family, member V (H2AFV), transcript variant 2, i
 NM_138636 Homo sapiens toll-like receptor 8 (TLR8), transcript variant 2, mRNA
 NM_138637 Homo sapiens dudulin 2 (TSAP6), mRNA
 NM_138638 Homo sapiens cofilin 2 (muscle) (CFL2), transcript variant 2, mRNA

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NM_138639 Homo sapiens BCL2-like 12 (proline rich) (BCL2L12), transcript variant 1, mf
 NM_138640 Homo sapiens golgi associated, gamma adaptin ear containing, ARF binding
 NM_138643 Homo sapiens calcium-binding tyrosine-(Y)-phosphorylation regulated (fibro.
 NM_138644 Homo sapiens calcium-binding tyrosine-(Y)-phosphorylation regulated (fibro.
 NM_138687 Homo sapiens phosphatidylinositol-4-phosphate 5-kinase, type II, beta (PIP5
 NM_138688 Homo sapiens toll-like receptor 9 (TLR9), transcript variant E, mRNA
 NM_138691 Homo sapiens transmembrane channel-like 1 (TMC1), mRNA
 NM_138693 Homo sapiens Kruppel-like factor 14 (KLF14), mRNA
 NM_138694 Homo sapiens polycystic kidney and hepatic disease 1 (autosomal recessive
 NM_138697 Homo sapiens taste receptor, type 1, member 1 (TAS1R1), transcript variant
 NM_138698 Homo sapiens prematurely terminated mRNA decay factor-like (LOC91431),
 NM_138699 Homo sapiens hypothetical protein BC006130 (LOC93622), mRNA
 NM_138700 Homo sapiens tripartite motif-containing 40 (TRIM40), mRNA
 NM_138701 Homo sapiens chromosome 7 open reading frame 11 (C7orf11), mRNA
 NM_138702 Homo sapiens melanoma antigen, family C, 3 (MAGEC3), transcript variant 1
 NM_138703 Homo sapiens melanoma antigen, family E, 2 (MAGEE2), mRNA
 NM_138704 Homo sapiens necdin-like 2 (NDNL2), mRNA
 NM_138705 Homo sapiens calgandulin-like protein (CAGLP), mRNA
 NM_138706 Homo sapiens beta-1,3-N-acetylglucosaminyltransferase protein (IMAGE49f
 NM_138707 Homo sapiens B-cell CLL/lymphoma 7B (BCL7B), transcript variant 2, mRNA
 NM_138709 Homo sapiens DAB2 Interacting protein (DAB2IP), mRNA
 NM_138711 Homo sapiens peroxisome proliferative activated receptor, gamma (PPARG)
 NM_138712 Homo sapiens peroxisome proliferative activated receptor, gamma (PPARG)
 NM_138713 Homo sapiens nuclear factor of activated T-cells 5, tonicity-responsive (NFA
 NM_138714 Homo sapiens nuclear factor of activated T-cells 5, tonicity-responsive (NFA
 NM_138715 Homo sapiens macrophage scavenger receptor 1 (MSR1), transcript variant
 NM_138716 Homo sapiens macrophage scavenger receptor 1 (MSR1), transcript variant
 NM_138717 Homo sapiens palmitoyl-protein thioesterase 2 (PPT2), transcript variant 2, n
 NM_138718 Homo sapiens solute carrier family 26, member 8 (SLC26A8), transcript varie
 NM_138720 Homo sapiens histone 1, H2bd (HIST1H2BD), transcript variant 2, mRNA
 NM_138722 Homo sapiens BCL2-like 14 (apoptosis facilitator) (BCL2L14), transcript vari
 NM_138723 Homo sapiens BCL2-like 14 (apoptosis facilitator) (BCL2L14), transcript vari
 NM_138724 Homo sapiens BCL2-like 14 (apoptosis facilitator) (BCL2L14), transcript vari
 NM_138726 Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 13
 NM_138727 Homo sapiens suppression of tumorigenicity 7 like (ST7L), transcript variant
 NM_138728 Homo sapiens suppression of tumorigenicity 7 like (ST7L), transcript variant
 NM_138729 Homo sapiens suppression of tumorigenicity 7 like (ST7L), transcript variant
 NM_138730 Homo sapiens high mobility group nucleosome binding domain 3 (HMGN3),
 NM_138731 Homo sapiens mirror-image polydactyly 1 (MIPOL1), mRNA
 NM_138732 Homo sapiens neurexin 2 (NRXN2), transcript variant alpha-2, mRNA
 NM_138733 Homo sapiens phosphoglycerate kinase 2 (PGK2), mRNA
 NM_138734 Homo sapiens neurexin 2 (NRXN2), transcript variant beta, mRNA
 NM_138735 Homo sapiens neurexin 1 (NRXN1), transcript variant beta, mRNA
 NM_138736 Homo sapiens guanine nucleotide binding protein (G protein), alpha activatin
 NM_138737 Homo sapiens hephaestin (HEPH), transcript variant 1, mRNA
 NM_138738 Homo sapiens SH2 domain containing phosphatase anchor protein 1 (SPAP
 NM_138739 Homo sapiens SH2 domain containing phosphatase anchor protein 1 (SPAP
 NM_138740 Homo sapiens NICE-3 protein (NICE-3), mRNA
 NM_138761 Homo sapiens BCL2-associated X protein (BAX), transcript variant alpha, mf
 NM_138762 Homo sapiens BCL2-associated X protein (BAX), transcript variant gamma, r
 NM_138763 Homo sapiens BCL2-associated X protein (BAX), transcript variant delta, mR
 NM_138764 Homo sapiens BCL2-associated X protein (BAX), transcript variant epsilon, n
 NM_138765 Homo sapiens BCL2-associated X protein (BAX), transcript variant sigma, m
 NM_138766 Homo sapiens peptidylglycine alpha-amidating monooxygenase (PAM), trans
 NM_138768 Homo sapiens myeloma overexpressed gene (in a subset of t(11;14) positive
 NM_138769 Homo sapiens ras homolog gene family, member T2 (RHOT2), mRNA
 NM_138770 Homo sapiens hypothetical protein BC016861 (LOC90557), mRNA

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NM_138771 Homo sapiens alpha-1,3(6)-mannosylglycoprotein beta-1,6-N-acetyl-glucosa
 NM_138773 Homo sapiens hypothetical protein BC017169 (LOC91137), mRNA
 NM_138774 Homo sapiens chromosome 19 open reading frame 22 (C19orf22), mRNA
 NM_138775 Homo sapiens hypothetical protein BC015183 (LOC91801), mRNA
 NM_138777 Homo sapiens mitochondrial ribosome recycling factor (MRRF), transcript va
 NM_138778 Homo sapiens chromosome 9 open reading frame 112 (C9orf112), mRNA
 NM_138779 Homo sapiens hypothetical protein BC015148 (LOC93081), mRNA
 NM_138780 Homo sapiens synaptotagmin-like 5 (SYTL5), mRNA
 NM_138781 Homo sapiens similar to envelope protein (LOC113386), mRNA
 NM_138783 Homo sapiens zinc finger protein Zip67 (ZIP67), mRNA
 NM_138784 Homo sapiens hypothetical protein BC014341 (LOC116123), mRNA
 NM_138785 Homo sapiens chromosome 6 open reading frame 72 (C6orf72), mRNA
 NM_138786 Homo sapiens hypothetical protein BC014339 (LOC116441), mRNA
 NM_138787 Homo sapiens hypothetical protein BC009561 (LOC119710), mRNA
 NM_138788 Homo sapiens hypothetical protein BC016153 (LOC120224), mRNA
 NM_138789 Homo sapiens hypothetical protein BC019238 (LOC120379), mRNA
 NM_138790 Homo sapiens hypothetical protein BC015003 (LOC122618), mRNA
 NM_138791 Homo sapiens chromosome 14 open reading frame 148 (C14orf148), mRNA
 NM_138792 Homo sapiens senescence downregulated lecl-like (LOC123169), mRNA
 NM_138793 Homo sapiens ectonucleoside triphosphate diphosphohydrolase 8 (ENTPD8)
 NM_138794 Homo sapiens lysophospholipase-like 1 (LYPLAL1), mRNA
 NM_138795 Homo sapiens ADP-ribosylation factor-like 10B (ARL10B), mRNA
 NM_138796 Homo sapiens hypothetical protein BC014608 (LOC128153), mRNA
 NM_138797 Homo sapiens hypothetical protein BC014641 (LOC129138), mRNA
 NM_138798 Homo sapiens hypothetical protein BC018453 (LOC129531), mRNA
 NM_138799 Homo sapiens O-acyltransferase (membrane bound) domain containing 2 (O
 NM_138800 Homo sapiens tripartite motif-containing 43 (TRIM43), mRNA
 NM_138801 Homo sapiens galactose mutarotase (aldose 1-epimerase) (GALM), mRNA
 NM_138802 Homo sapiens hypothetical protein BC018415 (LOC130617), mRNA
 NM_138803 Homo sapiens hypothetical protein BC015395 (LOC130940), mRNA
 NM_138804 Homo sapiens hypothetical protein BC014602 (LOC130951), mRNA
 NM_138805 Homo sapiens family with sequence similarity 3, member D (FAM3D), mRNA
 NM_138806 Homo sapiens MOX2 receptor (MOX2R), transcript variant 1, mRNA
 NM_138807 Homo sapiens hypothetical protein BC015088 (MGC16471), mRNA
 NM_138808 Homo sapiens hypothetical protein BC015210 (LOC132200), mRNA
 NM_138809 Homo sapiens hypothetical protein BC001573 (LOC134147), mRNA
 NM_138810 Homo sapiens T-cell activation GTPase activating protein (TAGAP), transcrip
 NM_138811 Homo sapiens chromosome 7 open reading frame 31 (C7orf31), mRNA
 NM_138812 Homo sapiens hypothetical protein BC019250 (LOC143241), mRNA
 NM_138813 Homo sapiens ATPase, Class I, type 8B, member 3 (ATP8B3), mRNA
 NM_138814 Homo sapiens GS2 like (LOC150379), mRNA
 NM_138815 Homo sapiens hypothetical protein BC018070 (LOC151871), mRNA
 NM_138817 Homo sapiens solute carrier family 7, (cationic amino acid transporter, y+ sy
 NM_138818 Homo sapiens chromosome 9 open reading frame 65 (C9orf65), mRNA
 NM_138819 Homo sapiens hypothetical protein BC017868 (LOC159091), mRNA
 NM_138820 Homo sapiens hypothetical protein MGC2198 (MGC2198), mRNA
 NM_138821 Homo sapiens peptidylglycine alpha-amidating monooxygenase (PAM), trans
 NM_138822 Homo sapiens peptidylglycine alpha-amidating monooxygenase (PAM), trans
 NM_138923 Homo sapiens TAF1 RNA polymerase II, TATA box binding protein (TBP)-as
 NM_138924 Homo sapiens guanidinacetate N-methyltransferase (GAMT), transcript vari
 NM_138925 Homo sapiens SON DNA binding protein (SON), transcript variant a, mRNA
 NM_138926 Homo sapiens SON DNA binding protein (SON), transcript variant c, mRNA
 NM_138927 Homo sapiens SON DNA binding protein (SON), transcript variant f, mRNA
 NM_138928 Homo sapiens molybdenum cofactor synthesis 1 (MOCS1), transcript variant
 NM_138929 Homo sapiens diablo homolog (Drosophila) (DIABLO), nuclear gene encodin
 NM_138930 Homo sapiens diablo homolog (Drosophila) (DIABLO), nuclear gene encodin
 NM_138931 Homo sapiens B-cell CLL/lymphoma 6 (zinc finger protein 51) (BCL6), transc

NM_138932 Homo sapiens apobec-1 complementation factor (ACF), transcript variant 2,
 NM_138933 Homo sapiens apobec-1 complementation factor (ACF), transcript variant 3,
 NM_138934 Homo sapiens palmitoyl-protein thioesterase 2 (PPT2), transcript variant 3, n
 NM_138937 Homo sapiens pancreatitis-associated protein (PAP), transcript variant 3, mF
 NM_138938 Homo sapiens pancreatitis-associated protein (PAP), transcript variant 2, mF
 NM_138939 Homo sapiens MOX2 receptor (MOX2R), transcript variant 2, mRNA
 NM_138940 Homo sapiens MOX2 receptor (MOX2R), transcript variant 3, mRNA
 NM_138957 Homo sapiens mitogen-activated protein kinase 1 (MAPK1), transcript varian
 NM_138958 Homo sapiens autocrine motility factor receptor (AMFR), transcript variant 2,
 NM_138959 Homo sapiens vang-like 1 (van gogh, Drosophila) (VANGL1), mRNA
 NM_138960 Homo sapiens TGF β -induced factor 2-like, X-linked (TGIF2LX), mRNA
 NM_138961 Homo sapiens endothelial cell adhesion molecule (ESAM), mRNA
 NM_138962 Homo sapiens musashi homolog 2 (Drosophila) (MSI2), transcript variant 1, i
 NM_138963 Homo sapiens ribosomal protein S4, Y-linked 2 (RPS4Y2), mRNA
 NM_138964 Homo sapiens G protein-coupled receptor 73 (GPR73), mRNA
 NM_138966 Homo sapiens neuropilin (NRP) and tolloid (TLL)-like 1 (NETO1), transcript v
 NM_138967 Homo sapiens secretory carrier membrane protein 5 (SCAMP5), mRNA
 NM_138969 Homo sapiens retinal short chain dehydrogenase reductase (RDH-E2), mRN
 NM_138970 Homo sapiens neuroligin 3 (NRXN3), transcript variant beta, mRNA
 NM_138971 Homo sapiens beta-site APP-cleaving enzyme 1 (BACE1), transcript variant
 NM_138972 Homo sapiens beta-site APP-cleaving enzyme 1 (BACE1), transcript variant
 NM_138973 Homo sapiens beta-site APP-cleaving enzyme 1 (BACE1), transcript variant
 NM_138980 Homo sapiens mitogen-activated protein kinase 10 (MAPK10), transcript vari
 NM_138981 Homo sapiens mitogen-activated protein kinase 10 (MAPK10), transcript vari
 NM_138982 Homo sapiens mitogen-activated protein kinase 10 (MAPK10), transcript vari
 NM_138983 Homo sapiens oligodendrocyte transcription factor 1 (OLIG1), mRNA
 NM_138991 Homo sapiens beta-site APP-cleaving enzyme 2 (BACE2), transcript variant
 NM_138992 Homo sapiens beta-site APP-cleaving enzyme 2 (BACE2), transcript variant
 NM_138993 Homo sapiens mitogen-activated protein kinase 11 (MAPK11), transcript vari
 NM_138994 Homo sapiens contactin associated protein-like 4 (CNTNAP4), transcript vari
 NM_138995 Homo sapiens myosin IIIB (MYO3B), mRNA
 NM_138996 Homo sapiens contactin associated protein-like 5 (CNTNAP5), transcript vari
 NM_138998 Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 39 (DDX39), transc
 NM_138999 Homo sapiens neuropilin (NRP) and tolloid (TLL)-like 1 (NETO1), transcript v
 NM_139002 Homo sapiens hemochromatosis (HFE), transcript variant 2, mRNA
 NM_139003 Homo sapiens hemochromatosis (HFE), transcript variant 3, mRNA
 NM_139004 Homo sapiens hemochromatosis (HFE), transcript variant 4, mRNA
 NM_139005 Homo sapiens hemochromatosis (HFE), transcript variant 5, mRNA
 NM_139006 Homo sapiens hemochromatosis (HFE), transcript variant 6, mRNA
 NM_139007 Homo sapiens hemochromatosis (HFE), transcript variant 7, mRNA
 NM_139008 Homo sapiens hemochromatosis (HFE), transcript variant 8, mRNA
 NM_139009 Homo sapiens hemochromatosis (HFE), transcript variant 9, mRNA
 NM_139010 Homo sapiens hemochromatosis (HFE), transcript variant 10, mRNA
 NM_139011 Homo sapiens hemochromatosis (HFE), transcript variant 11, mRNA
 NM_139012 Homo sapiens mitogen-activated protein kinase 14 (MAPK14), transcript vari
 NM_139013 Homo sapiens mitogen-activated protein kinase 14 (MAPK14), transcript vari
 NM_139014 Homo sapiens mitogen-activated protein kinase 14 (MAPK14), transcript vari
 NM_139015 Homo sapiens signal peptide peptidase 3 (SPPL3), mRNA
 NM_139016 Homo sapiens hypothetical gene LOC128439 (LOC128439), mRNA
 NM_139017 Homo sapiens interleukin 31 receptor A (IL31RA), mRNA
 NM_139018 Homo sapiens NK inhibitory receptor precursor (NKIR), mRNA
 NM_139021 Homo sapiens extracellular signal-regulated kinase 8 (ERK8), mRNA
 NM_139022 Homo sapiens pan-hematopoietic expression (PHEMX), transcript variant 1,
 NM_139024 Homo sapiens pan-hematopoietic expression (PHEMX), transcript variant 3,
 NM_139025 Homo sapiens a disintegrin-like and metalloprotease (repolydin type) with th
 NM_139026 Homo sapiens a disintegrin-like and metalloprotease (repolydin type) with th
 NM_139027 Homo sapiens a disintegrin-like and metalloprotease (repolydin type) with th

NM_139028 Homo sapiens a disintegrin-like and metalloprotease (repolysin type) with th
 NM_139030 Homo sapiens CD151 antigen (CD151), transcript variant 2, mRNA
 NM_139032 Homo sapiens mitogen-activated protein kinase 7 (MAPK7), transcript varian
 NM_139033 Homo sapiens mitogen-activated protein kinase 7 (MAPK7), transcript varian
 NM_139034 Homo sapiens mitogen-activated protein kinase 7 (MAPK7), transcript varian
 NM_139035 Homo sapiens SWI/SNF related, matrix associated, actin dependent regulat
 NM_139045 Homo sapiens SWI/SNF related, matrix associated, actin dependent regulat
 NM_139046 Homo sapiens mitogen-activated protein kinase 8 (MAPK8), transcript varian
 NM_139047 Homo sapiens mitogen-activated protein kinase 8 (MAPK8), transcript varian
 NM_139048 Homo sapiens SWI/SNF related, matrix associated, actin dependent regulat
 NM_139049 Homo sapiens mitogen-activated protein kinase 8 (MAPK8), transcript varian
 NM_139053 Homo sapiens EPS8-like 3 (EPS8L3), transcript variant 1, mRNA
 NM_139054 Homo sapiens a disintegrin-like and metalloprotease (repolysin type) with th
 NM_139055 Homo sapiens a disintegrin-like and metalloprotease (repolysin type) with th
 NM_139056 Homo sapiens a disintegrin-like and metalloprotease (repolysin type) with th
 NM_139057 Homo sapiens a disintegrin-like and metalloprotease (repolysin type) with th
 NM_139058 Homo sapiens aristaleless related homeobox (ARX), mRNA
 NM_139062 Homo sapiens casein kinase 1, delta (CSNK1D), transcript variant 2, mRNA
 NM_139067 Homo sapiens SWI/SNF related, matrix associated, actin dependent regulat
 NM_139068 Homo sapiens mitogen-activated protein kinase 9 (MAPK9), transcript varian
 NM_139069 Homo sapiens mitogen-activated protein kinase 9 (MAPK9), transcript varian
 NM_139070 Homo sapiens mitogen-activated protein kinase 9 (MAPK9), transcript varian
 NM_139071 Homo sapiens SWI/SNF related, matrix associated, actin dependent regulat
 NM_139072 Homo sapiens delta-notch-like EGF repeat-containing transmembrane (DNE
 NM_139073 Homo sapiens spermatogenesis associated 3 (SPATA3), mRNA
 NM_139074 Homo sapiens defensin, beta 127 (DEFB127), mRNA
 NM_139075 Homo sapiens two pore segment channel 2 (TPCN2), mRNA
 NM_139076 Homo sapiens hypothetical protein FLJ13614 (FLJ13614), mRNA
 NM_139078 Homo sapiens mitogen-activated protein kinase-activated protein kinase 5 (l
 NM_139118 Homo sapiens YY1 associated protein (YAP), transcript variant 2, mRNA
 NM_139119 Homo sapiens YY1 associated protein (YAP), transcript variant 3, mRNA
 NM_139120 Homo sapiens YY1 associated protein (YAP), transcript variant 4, mRNA
 NM_139121 Homo sapiens YY1 associated protein (YAP), transcript variant 5, mRNA
 NM_139122 Homo sapiens TAF6 RNA polymerase II, TATA box binding protein (TBP)-as
 NM_139123 Homo sapiens TAF6 RNA polymerase II, TATA box binding protein (TBP)-as
 NM_139124 Homo sapiens mitogen-activated protein kinase 8 interacting protein 2 (MAPI
 NM_139125 Homo sapiens mannan-binding lectin serine protease 1 (C4/C2 activating coi
 NM_139126 Homo sapiens peptidylprolyl isomerase (cyclophilin)-like 4 (PPIL4), mRNA
 NM_139131 Homo sapiens nucleoporin 98kDa (NUP98), transcript variant 2, mRNA
 NM_139132 Homo sapiens nucleoporin 98kDa (NUP98), transcript variant 4, mRNA
 NM_139135 Homo sapiens AT rich interactive domain 1A (SWI- like) (ARID1A), transcript
 NM_139136 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, m
 NM_139137 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, m
 NM_139155 Homo sapiens a disintegrin-like and metalloprotease (repolysin type) with th
 NM_139156 Homo sapiens adenosine monophosphate deaminase 2 (isoform L) (AMPD2
 NM_139157 Homo sapiens suppression of tumorigenicity 5 (ST5), transcript variant 2, mF
 NM_139158 Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region,
 NM_139159 Homo sapiens dipeptidylpeptidase 9 (DPP9), mRNA
 NM_139160 Homo sapiens novel 58.3 KDA protein (LOC91614), mRNA
 NM_139161 Homo sapiens crumbs homolog 3 (Drosophila) (CRB3), transcript variant 2, r
 NM_139162 Homo sapiens Smith-Magenis syndrome chromosome region, candidate 7 (S
 NM_139163 Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region,
 NM_139164 Homo sapiens START domain containing 4, sterol regulated (STARD4), mR1
 NM_139165 Homo sapiens retinoic acid early transcript 1E (RAET1E), mRNA
 NM_139166 Homo sapiens striated muscle activator of Rho-dependent signaling (STARS
 NM_139167 Homo sapiens sarcoglycan zeta (SGCZ), mRNA
 NM_139168 Homo sapiens splicing factor, arginine/serine-rich 12 (SFRS12), mRNA

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NM_139169 Homo sapiens TruB pseudouridine (psi) synthase homolog 1 (E. coli) (TRUB
 NM_139170 Homo sapiens hypothetical protein AF447587 (LOC146562), mRNA
 NM_139171 Homo sapiens START domain containing 6 (STARD6), mRNA
 NM_139172 Homo sapiens MDAC1 (MDAC1), mRNA
 NM_139173 Homo sapiens CG10806-like (LOC150159), mRNA
 NM_139174 Homo sapiens testis nuclear RNA-binding protein-like (LOC161931), mRNA
 NM_139175 Homo sapiens ring finger protein 133 (RNF133), mRNA
 NM_139176 Homo sapiens NACHT, leucine rich repeat and PYD containing 7 (NALP7), t
 NM_139177 Homo sapiens solute carrier family 39 (metal ion transporter), member 11 (S
 NM_139178 Homo sapiens prostate cancer antigen-1 (DEPC-1), mRNA
 NM_139179 Homo sapiens KCCR13L (LOC221955), mRNA
 NM_139181 Homo sapiens centaurin, delta 2 (CENTD2), transcript variant 1, mRNA
 NM_139182 Homo sapiens centaurin, delta 1 (CENTD1), transcript variant 2, mRNA
 NM_139199 Homo sapiens bromodomain containing 8 (BRD8), transcript variant 2, mRN
 NM_139201 Homo sapiens G protein-coupled receptor kinase interactor 2 (GIT2), transcr
 NM_139202 Homo sapiens megalencephalic leukoencephalopathy with subcortical cysts
 NM_139204 Homo sapiens EPS8-like 1 (EPS8L1), transcript variant 3, mRNA
 NM_139205 Homo sapiens histone deacetylase 5 (HDAC5), transcript variant 2, mRNA
 NM_139207 Homo sapiens nucleosome assembly protein 1-like 1 (NAP1L1), transcript ve
 NM_139208 Homo sapiens mannan-binding lectin serine protease 2 (MASP2), transcript
 NM_139209 Homo sapiens G protein-coupled receptor kinase 7 (GRK7), mRNA
 NM_139211 Homo sapiens homeodomain-only protein (HOP), transcript variant 2, mRNA
 NM_139212 Homo sapiens homeodomain-only protein (HOP), transcript variant 3, mRNA
 NM_139214 Homo sapiens TGFB-induced factor 2-like, Y-linked (TGIF2LY), mRNA
 NM_139215 Homo sapiens TAF15 RNA polymerase II, TATA box binding protein (TBP)-a
 NM_139235 Homo sapiens nucleolar protein family 8 (RNA-associated) (NOL6), transcript
 NM_139238 Homo sapiens ADAMTS-like 1 (ADAMTSL1), transcript variant 1, mRNA
 NM_139239 Homo sapiens T-cell activation NFkB-like protein (TA-NFKBH), mRNA
 NM_139240 Homo sapiens LOC92346 (LOC92346), mRNA
 NM_139241 Homo sapiens FGD1 family, member 4 (FGD4), mRNA
 NM_139242 Homo sapiens methionyl-tRNA formyltransferase, mitochondrial (MFMFT), m
 NM_139243 Homo sapiens testis nuclear RNA-binding protein (Tenr), mRNA
 NM_139244 Homo sapiens syntaxin binding protein 5 (tomosyn) (STXBP5), mRNA
 NM_139245 Homo sapiens protein phosphatase 1 (formerly 2C)-like (PPM1L), mRNA
 NM_139246 Homo sapiens chromosome 9 open reading frame 97 (C9orf97), mRNA
 NM_139247 Homo sapiens adenylate cyclase 4 (ADCY4), mRNA
 NM_139248 Homo sapiens lipase, member H (LIPH), mRNA
 NM_139249 Homo sapiens membrane-spanning 4-domains, subfamily A, member 6E (M
 NM_139250 Homo sapiens cancer/testis antigen 1A (CTAG1A), mRNA
 NM_139264 Homo sapiens ADAMTS-like 1 (ADAMTSL1), transcript variant 3, mRNA
 NM_139265 Homo sapiens EH-domain containing 4 (EHD4), mRNA
 NM_139266 Homo sapiens signal transducer and activator of transcription 1, 91kDa (STA
 NM_139267 Homo sapiens START domain containing 7 (STARD7), transcript variant 2, n
 NM_139273 Homo sapiens cysteinyl-tRNA synthetase (CARS), transcript variant 1, mRN
 NM_139274 Homo sapiens acetyl-Coenzyme A synthetase 2 (ADP forming) (ACAS2), tra
 NM_139275 Homo sapiens A kinase (PRKA) anchor protein 1 (AKAP1), nuclear gene enc
 NM_139276 Homo sapiens signal transducer and activator of transcription 3 (acute-phase
 NM_139277 Homo sapiens kallikrein 7 (chymotryptic, stratum corneum) (KLK7), transcrip
 NM_139278 Homo sapiens leucine-rich repeat LGI family, member 3 (LGI3), mRNA
 NM_139279 Homo sapiens multiple coagulation factor deficiency 2 (MCFD2), mRNA
 NM_139280 Homo sapiens ORM1-like 3 (S. cerevisiae) (ORMDL3), mRNA
 NM_139281 Homo sapiens WD repeat domain 36 (WDR36), mRNA
 NM_139282 Homo sapiens paired-like homeobox protein OTEX (OTEX), mRNA
 NM_139283 Homo sapiens T-cell activation protein phosphatase 2C (TA-PP2C), mRNA
 NM_139284 Homo sapiens leucine-rich repeat LGI family, member 4 (LGI4), mRNA
 NM_139285 Homo sapiens growth arrest-specific 2 like 2 (GAS2L2), mRNA
 NM_139286 Homo sapiens cell division cycle 26 (CDC26), mRNA

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NM_139289 Homo sapiens A kinase (PRKA) anchor protein 4 (AKAP4), transcript variant
 NM_139290 Homo sapiens angiotensin II (ANGPT1), transcript variant 2, mRNA
 NM_139312 Homo sapiens YME1-like 1 (S. cerevisiae) (YME1L1), nuclear gene encoding
 NM_139313 Homo sapiens YME1-like 1 (S. cerevisiae) (YME1L1), nuclear gene encoding
 NM_139314 Homo sapiens angiotensin-like 4 (ANGPTL4), transcript variant 1, mRNA
 NM_139315 Homo sapiens TAF6 RNA polymerase II, TATA box binding protein (TBP)-as
 NM_139316 Homo sapiens amphiphysin (Stiff-Man syndrome with breast cancer 128kDa
 NM_139317 Homo sapiens baculoviral IAP repeat-containing 7 (livin) (BIRC7), transcript
 NM_139318 Homo sapiens potassium voltage-gated channel, subfamily H (eag-related), I
 NM_139319 Homo sapiens solute carrier family 17 (sodium-dependent inorganic phosph
 NM_139320 Homo sapiens CHRNA7 (cholinergic receptor, nicotinic, alpha polypeptide 7,
 NM_139321 Homo sapiens attractin (ATRIN), transcript variant 1, mRNA
 NM_139322 Homo sapiens attractin (ATRIN), transcript variant 2, mRNA
 NM_139323 Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase act
 NM_139343 Homo sapiens bridging integrator 1 (BIN1), transcript variant 1, mRNA
 NM_139344 Homo sapiens bridging integrator 1 (BIN1), transcript variant 2, mRNA
 NM_139345 Homo sapiens bridging integrator 1 (BIN1), transcript variant 3, mRNA
 NM_139346 Homo sapiens bridging integrator 1 (BIN1), transcript variant 4, mRNA
 NM_139347 Homo sapiens bridging integrator 1 (BIN1), transcript variant 5, mRNA
 NM_139348 Homo sapiens bridging integrator 1 (BIN1), transcript variant 6, mRNA
 NM_139349 Homo sapiens bridging integrator 1 (BIN1), transcript variant 7, mRNA
 NM_139350 Homo sapiens bridging integrator 1 (BIN1), transcript variant 8, mRNA
 NM_139351 Homo sapiens bridging integrator 1 (BIN1), transcript variant 10, mRNA
 NM_139352 Homo sapiens TATA box binding protein (TBP)-associated factor, RNA polym
 NM_139353 Homo sapiens TATA box binding protein (TBP)-associated factor, RNA polym
 NM_139354 Homo sapiens megakaryocyte-associated tyrosine kinase (MATK), transcript
 NM_139355 Homo sapiens megakaryocyte-associated tyrosine kinase (MATK), transcript
 NM_144488 Homo sapiens regulator of G-protein signalling 3 (RGS3), transcript variant 6
 NM_144489 Homo sapiens regulator of G-protein signalling 3 (RGS3), transcript variant 5
 NM_144490 Homo sapiens A kinase (PRKA) anchor protein 11 (AKAP11), transcript varie
 NM_144492 Homo sapiens claudin 14 (CLDN14), transcript variant 1, mRNA
 NM_144494 Homo sapiens polyglutamine binding protein 1 (PQBP1), mRNA
 NM_144495 Homo sapiens polyglutamine binding protein 1 (PQBP1), mRNA
 NM_144497 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), transc
 NM_144498 Homo sapiens oxysterol binding protein-like 2 (OSBPL2), transcript variant 2
 NM_144499 Homo sapiens guanine nucleotide binding protein (G protein), alpha transduc
 NM_144501 Homo sapiens F11 receptor (F11R), transcript variant 2, mRNA
 NM_144502 Homo sapiens F11 receptor (F11R), transcript variant 3, mRNA
 NM_144503 Homo sapiens F11 receptor (F11R), transcript variant 4, mRNA
 NM_144504 Homo sapiens F11 receptor (F11R), transcript variant 5, mRNA
 NM_144505 Homo sapiens kallikrein 8 (neuropsin/ovasin) (KLK8), transcript variant 2, mF
 NM_144506 Homo sapiens kallikrein 8 (neuropsin/ovasin) (KLK8), transcript variant 3, mF
 NM_144507 Homo sapiens kallikrein 8 (neuropsin/ovasin) (KLK8), transcript variant 4, mF
 NM_144508 Homo sapiens AF15Q14 protein (AF15Q14), mRNA
 NM_144563 Homo sapiens ribose 5-phosphate isomerase A (ribose 5-phosphate epimer
 NM_144564 Homo sapiens solute carrier family 39 (zinc transporter), member 3 (SLC39A
 NM_144565 Homo sapiens homolog of Drosophila Numb-interacting protein (NIP), mRNA
 NM_144567 Homo sapiens similar to RIKEN cDNA 2610307121 (LOC90806), mRNA
 NM_144568 Homo sapiens chromosome 14 open reading frame 9 (C14orf9), mRNA
 NM_144569 Homo sapiens hypothetical protein FLJ25348 (FLJ25348), mRNA
 NM_144570 Homo sapiens chromosome 16 open reading frame 34 (C16orf34), mRNA
 NM_144571 Homo sapiens CCR4-NOT transcription complex, subunit 6-like (CNOT6L), n
 NM_144573 Homo sapiens nexilin (F actin binding protein) (NEXN), mRNA
 NM_144574 Homo sapiens WD repeat domain 20 (WDR20), transcript variant 2, mRNA
 NM_144575 Homo sapiens calpain 13 (CAPN13), mRNA
 NM_144576 Homo sapiens hypothetical protein FLJ32452 (FLJ32452), mRNA
 NM_144577 Homo sapiens hypothetical protein FLJ32926 (FLJ32926), mRNA

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NM_144578 Homo sapiens chromosome 14 open reading frame 32 (C14orf32), mRNA
 NM_144579 Homo sapiens sideroflexin 5 (SFXN5), mRNA
 NM_144580 Homo sapiens kidney predominant protein NCU-G1 (MGC31963), mRNA
 NM_144581 Homo sapiens chromosome 14 open reading frame 149 (C14orf149), mRNA
 NM_144582 Homo sapiens testis expressed sequence 261 (TEX261), mRNA
 NM_144583 Homo sapiens ATPase, H+ transporting, lysosomal 42kDa, V1 subunit C isoform 1 (ATP13C1), mRNA
 NM_144584 Homo sapiens hypothetical protein FLJ30525 (FLJ30525), mRNA
 NM_144585 Homo sapiens solute carrier family 22 (organic anion/cation transporter), member 1 (SLC22A1), mRNA
 NM_144586 Homo sapiens hypothetical protein MGC29643 (MGC29643), mRNA
 NM_144587 Homo sapiens chromosome 10 open reading frame 87 (C10orf87), mRNA
 NM_144588 Homo sapiens zinc finger, FYVE domain containing 27 (ZFYVE27), transcript variant 1, mRNA
 NM_144589 Homo sapiens catechol-O-methyltransferase domain containing 1 (COMT D1), mRNA
 NM_144590 Homo sapiens ankyrin repeat domain 22 (ANKRD22), mRNA
 NM_144591 Homo sapiens chromosome 10 open reading frame 32 (C10orf32), mRNA
 NM_144593 Homo sapiens Ras homolog enriched in brain like 1 (RHEBL1), mRNA
 NM_144594 Homo sapiens hypothetical protein FLJ32942 (FLJ32942), mRNA
 NM_144595 Homo sapiens hypothetical protein FLJ30046 (FLJ30046), mRNA
 NM_144596 Homo sapiens tetratricopeptide repeat domain 8 (TTC8), transcript variant 3, mRNA
 NM_144597 Homo sapiens hypothetical protein MGC29937 (MGC29937), mRNA
 NM_144598 Homo sapiens leucine rich repeat containing 28 (LRRC28), mRNA
 NM_144599 Homo sapiens non-imprinted in Prader-Willi/Angelman syndrome 1 (NIPA1), mRNA
 NM_144600 Homo sapiens hypothetical protein FLJ31153 (FLJ31153), mRNA
 NM_144601 Homo sapiens chemokine-like factor super family 3 (CKLFSF3), transcript variant 1, mRNA
 NM_144602 Homo sapiens hypothetical protein MGC32905 (MGC32905), mRNA
 NM_144603 Homo sapiens NADPH oxidase organizer 1 (NOXO1), transcript variant 1, mRNA
 NM_144604 Homo sapiens hypothetical protein BC001584 (LOC124245), mRNA
 NM_144605 Homo sapiens hypothetical protein FLJ25410 (FLJ25410), mRNA
 NM_144606 Homo sapiens folliculin (BHD), transcript variant 2, mRNA
 NM_144607 Homo sapiens hypothetical protein FLJ32499 (FLJ32499), mRNA
 NM_144608 Homo sapiens hypothetical protein MGC39389 (FLJ32384), mRNA
 NM_144609 Homo sapiens hypothetical protein FLJ31795 (FLJ31795), mRNA
 NM_144610 Homo sapiens hypothetical protein FLJ25006 (FLJ25006), mRNA
 NM_144611 Homo sapiens hypothetical protein MGC32124 (MGC32124), mRNA
 NM_144612 Homo sapiens lipoxygenase homology domains 1 (LOXHD1), mRNA
 NM_144613 Homo sapiens cytochrome c oxidase subunit Vlb, testes-specific (COXVB2), mRNA
 NM_144614 Homo sapiens methyl-CpG binding domain protein 3-like 2 (MBD3L2), mRNA
 NM_144615 Homo sapiens hypothetical protein MGC23244 (MGC23244), mRNA
 NM_144616 Homo sapiens homolog of mouse skeletal muscle sarcoplasmic reticulum protein 1 (HSPB6), mRNA
 NM_144617 Homo sapiens heat shock protein, alpha-crystallin-related, B6 (HSPB6), mRNA
 NM_144618 Homo sapiens hypothetical protein MGC29891 (MGC29891), mRNA
 NM_144620 Homo sapiens hypothetical protein MGC14816 (MGC14816), mRNA
 NM_144621 Homo sapiens zinc finger and BTB domain containing 8 (ZBTB8), mRNA
 NM_144622 Homo sapiens hypothetical protein FLJ32934 (FLJ32934), mRNA
 NM_144623 Homo sapiens hypothetical protein FLJ32784 (FLJ32784), mRNA
 NM_144624 Homo sapiens kinase interacting with leukemia-associated gene (statmin) (KIF18A), mRNA
 NM_144625 Homo sapiens hypothetical protein FLJ32978 (FLJ32978), mRNA
 NM_144626 Homo sapiens hypothetical protein MGC17299 (MGC17299), mRNA
 NM_144627 Homo sapiens SSTR-interacting protein (SSTR-IP), mRNA
 NM_144628 Homo sapiens chromosome 20 open reading frame 140 (C20orf140), mRNA
 NM_144629 Homo sapiens chromosome 2 open reading frame 11 (C2orf11), mRNA
 NM_144631 Homo sapiens zinc finger protein 513 (ZNF513), mRNA
 NM_144632 Homo sapiens hypothetical protein FLJ30294 (FLJ30294), mRNA
 NM_144633 Homo sapiens potassium voltage-gated channel, subfamily H (eag-related), member 1 (KCNH1), mRNA
 NM_144634 Homo sapiens lysozyme-like 4 (LYZL4), mRNA
 NM_144635 Homo sapiens hypothetical protein MGC21688 (MGC21688), mRNA
 NM_144636 Homo sapiens coiled-coil-helix-coiled-coil-helix domain containing 4 (CHCHC1), mRNA
 NM_144637 Homo sapiens zinc finger, DHHC domain containing 19 (ZDHHC19), mRNA

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NM_144638 Homo sapiens hypothetical protein MGC29956 (MGC29956), mRNA
NM_144639 Homo sapiens hypothetical protein FLJ31300 (FLJ31300), mRNA
NM_144640 Homo sapiens interleukin 17 receptor E (IL17RE), transcript variant 3, mRNA
NM_144641 Homo sapiens likely ortholog of mouse protein phosphatase 2C eta (FLJ323:
NM_144642 Homo sapiens synaptoporin (SYNPR), mRNA
NM_144643 Homo sapiens hypothetical protein FLJ30655 (FLJ30655), mRNA
NM_144644 Homo sapiens spermatogenesis associated 4 (SPATA4), mRNA
NM_144645 Homo sapiens hypothetical protein MGC26744 (MGC26744), mRNA
NM_144646 Homo sapiens immunoglobulin J polypeptide, linker protein for immunoglob
NM_144647 Homo sapiens hypothetical protein MGC26610 (MGC26610), mRNA
NM_144648 Homo sapiens hypothetical protein FLJ32786 (FLJ32786), mRNA
NM_144649 Homo sapiens hypothetical protein FLJ33069 (FLJ33069), mRNA
NM_144650 Homo sapiens alcohol dehydrogenase, iron containing, 1 (ADHFE1), mRNA
NM_144651 Homo sapiens hypothetical protein FLJ25471 (FLJ25471), mRNA
NM_144652 Homo sapiens leucine zipper-EF-hand containing transmembrane protein 2 (
NM_144653 Homo sapiens BTB (POZ) domain containing 14A (BTBD14A), mRNA
NM_144654 Homo sapiens chromosome 9 open reading frame 116 (C9orf116), mRNA
NM_144657 Homo sapiens hypothetical protein FLJ30678 (FLJ30678), mRNA
NM_144658 Homo sapiens dedicator of cytokinesis 11 (DOCK11), mRNA
NM_144659 Homo sapiens t-complex 10 (mouse)-like (TCP10L), mRNA
NM_144660 Homo sapiens sterile alpha motif domain containing 8 (SAMD8), mRNA
NM_144661 Homo sapiens chromosome 10 open reading frame 82 (C10orf82), mRNA
NM_144662 Homo sapiens hypothetical protein MGC26605 (MGC26605), mRNA
NM_144663 Homo sapiens NOV1 (NOV1), mRNA
NM_144664 Homo sapiens hypothetical protein MGC33371 (MGC33371), mRNA
NM_144665 Homo sapiens sestrin 3 (SESN3), mRNA
NM_144666 Homo sapiens hypothetical protein FLJ32752 (FLJ32752), mRNA
NM_144667 Homo sapiens hypothetical protein FLJ32894 (FLJ32894), mRNA
NM_144668 Homo sapiens hypothetical protein MGC33630 (MGC33630), mRNA
NM_144669 Homo sapiens hypothetical protein FLJ31978 (FLJ31978), mRNA
NM_144670 Homo sapiens hypothetical protein FLJ25179 (FLJ25179), mRNA
NM_144671 Homo sapiens hypothetical protein FLJ32356 (FLJ32356), mRNA
NM_144672 Homo sapiens otoacortin (OTOA), mRNA
NM_144673 Homo sapiens chemokine-like factor super family 2 (CKLFSF2), mRNA
NM_144674 Homo sapiens hypothetical protein FLJ32871 (FLJ32871), mRNA
NM_144675 Homo sapiens hypothetical protein MGC18079 (MGC18079), mRNA
NM_144676 Homo sapiens hypothetical protein MGC23911 (MGC23911), mRNA
NM_144677 Homo sapiens mannosyl (alpha-1,6)-glycoprotein beta-1,6-N-acetyl-glucosa
NM_144678 Homo sapiens target of myb1-like 2 (chicken) (TOM1L2), mRNA
NM_144679 Homo sapiens hypothetical protein FLJ31528 (FLJ31528), mRNA
NM_144681 Homo sapiens hypothetical protein FLJ32734 (FLJ32734), mRNA
NM_144682 Homo sapiens hypothetical protein FLJ31952 (FLJ31952), mRNA
NM_144683 Homo sapiens hypothetical protein MGC23280 (MGC23280), mRNA
NM_144684 Homo sapiens zinc finger protein 480 (ZNF480), mRNA
NM_144685 Homo sapiens homeodomain interacting protein kinase 4 (HIPK4), mRNA
NM_144686 Homo sapiens transmembrane channel-like 4 (TMC4), mRNA
NM_144687 Homo sapiens NACHT, leucine rich repeat and PYD containing 12 (NALP12)
NM_144688 Homo sapiens hypothetical protein FLJ32658 (FLJ32658), mRNA
NM_144689 Homo sapiens hypothetical protein FLJ32191 (FLJ32191), mRNA
NM_144690 Homo sapiens zinc finger protein 582 (ZNF582), mRNA
NM_144691 Homo sapiens calpain 12 (CAPN12), mRNA
NM_144692 Homo sapiens hypothetical protein BC017947 (LOC148137), mRNA
NM_144693 Homo sapiens zinc finger protein 558 (ZNF558), mRNA
NM_144694 Homo sapiens zinc finger protein 570 (ZNF570), mRNA
NM_144695 Homo sapiens hypothetical protein FLJ32421 (FLJ32421), mRNA
NM_144696 Homo sapiens hypothetical protein FLJ32940 (DKFZp686H1423), transcript
NM_144697 Homo sapiens hypothetical protein BC017397 (LOC148523), mRNA

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NM_144698 Homo sapiens hypothetical protein FLJ25124 (FLJ25124), mRNA
 NM_144699 Homo sapiens ATPase, Na⁺/K⁺ transporting, alpha 4 polypeptide (ATP1A4),
 NM_144701 Homo sapiens interleukin-23 receptor (IL23R), mRNA
 NM_144702 Homo sapiens hypothetical protein FLJ32884 (FLJ32884), mRNA
 NM_144703 Homo sapiens chromosome 20 open reading frame 40 (C20orf40), mRNA
 NM_144704 Homo sapiens hypothetical protein FLJ30473 (FLJ30473), mRNA
 NM_144705 Homo sapiens hypothetical protein MGC27019 (MGC27019), mRNA
 NM_144706 Homo sapiens chromosome 2 open reading frame 15 (C2orf15), mRNA
 NM_144707 Homo sapiens prominin 2 (PROM2), mRNA
 NM_144709 Homo sapiens hypothetical protein FLJ32312 (FLJ32312), mRNA
 NM_144710 Homo sapiens septin 10 (SEPT10), transcript variant 1, mRNA
 NM_144711 Homo sapiens hypothetical protein MGC22679 (MGC22679), mRNA
 NM_144712 Homo sapiens solute carrier family 23 (nucleobase transporters), member 3,
 NM_144713 Homo sapiens hypothetical protein FLJ32954 (FLJ32954), mRNA
 NM_144714 Homo sapiens hypothetical protein MGC27069 (FLJ25449), mRNA
 NM_144715 Homo sapiens hypothetical protein FLJ25200 (FLJ25200), mRNA
 NM_144716 Homo sapiens hypothetical protein MGC23918 (MGC23918), mRNA
 NM_144717 Homo sapiens hypothetical protein MGC34923 (MGC34923), mRNA
 NM_144718 Homo sapiens hypothetical protein AY099107 (LOC152185), mRNA
 NM_144719 Homo sapiens hypothetical protein FLJ25467 (FLJ25467), mRNA
 NM_144720 Homo sapiens multiple coiled-coil GABABR1-binding protein (MARLIN1), mF
 NM_144721 Homo sapiens THAP domain containing 6 (THAP6), mRNA
 NM_144722 Homo sapiens KPL2 protein (FLJ23577), transcript variant 2, mRNA
 NM_144723 Homo sapiens hypothetical protein FLJ31121 (FLJ31121), mRNA
 NM_144724 Homo sapiens MARVEL domain containing 2 (MARVELD2), mRNA
 NM_144725 Homo sapiens hypothetical protein FLJ25439 (FLJ25439), mRNA
 NM_144726 Homo sapiens hypothetical protein FLJ31951 (FLJ31951), mRNA
 NM_144727 Homo sapiens crystallin, gamma N (CRYGN), mRNA
 NM_144728 Homo sapiens dual specificity phosphatase 10 (DUSP10), transcript variant;
 NM_144729 Homo sapiens dual specificity phosphatase 10 (DUSP10), transcript variant;
 NM_144732 Homo sapiens heterogeneous nuclear ribonucleoprotein U-like 1 (HNRPUL1)
 NM_144733 Homo sapiens heterogeneous nuclear ribonucleoprotein U-like 1 (HNRPUL1)
 NM_144734 Homo sapiens heterogeneous nuclear ribonucleoprotein U-like 1 (HNRPUL1)
 NM_144738 Homo sapiens hypothetical protein PRO1853 (PRO1853), transcript variant 1
 NM_144765 Homo sapiens epithelial V-like antigen 1 (EVA1), transcript variant 2, mRNA
 NM_144768 Homo sapiens regulator of G-protein signalling 13 (RGS13), transcript varian
 NM_144767 Homo sapiens A kinase (PRKA) anchor protein 13 (AKAP13), transcript varie
 NM_144789 Homo sapiens forkhead box I1 (FOXI1), transcript variant 2, mRNA
 NM_144770 Homo sapiens RNA binding motif protein 11 (RBM11), mRNA
 NM_144772 Homo sapiens apolipoprotein A-I binding protein (APOA1BP), mRNA
 NM_144773 Homo sapiens G protein-coupled receptor 73-like 1 (GPR73L1), mRNA
 NM_144775 Homo sapiens Smith-Magenis syndrome chromosome region, candidate 8 (S
 NM_144776 Homo sapiens formyltetrahydrofolate dehydrogenase (FTHFD), transcript vai
 NM_144777 Homo sapiens scellin (SCEL), transcript variant 2, mRNA
 NM_144778 Homo sapiens muscleblind-like 2 (Drosophila) (MBNL2), transcript variant 1,
 NM_144779 Homo sapiens FXYD domain containing ion transport regulator 5 (FXYD5), tr
 NM_144780 Homo sapiens degenerative spermatocyte homolog, lipid desaturase (Droso
 NM_144781 Homo sapiens programmed cell death 2 (PDCD2), transcript variant 2, mRN
 NM_144782 Homo sapiens carnitine acetyltransferase (CRAT), transcript variant 3, mRN
 NM_144947 Homo sapiens kallikrein 11 (KLK11), transcript variant 2, mRNA
 NM_144949 Homo sapiens suppressor of cytokine signaling 5 (SOCS5), transcript variant
 NM_144956 Homo sapiens protease, serine, 21 (testisin) (PRSS21), transcript variant 2, i
 NM_144957 Homo sapiens protease, serine, 21 (testisin) (PRSS21), transcript variant 3, i
 NM_144962 Homo sapiens hypothetical protein MGC22776 (MGC22776), mRNA
 NM_144963 Homo sapiens hypothetical protein FLJ23790 (FLJ23790), mRNA
 NM_144964 Homo sapiens RNA (guanine-9-) methyltransferase domain containing 3 (RC
 NM_144965 Homo sapiens tetratricopeptide repeat domain 16 (TTC16), mRNA

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NM_144966 Homo sapiens chromosome 9 open reading frame 154 (C9orf154), mRNA
 NM_144967 Homo sapiens hypothetical protein FLJ30058 (FLJ30058), mRNA
 NM_144968 Homo sapiens hypothetical protein FLJ32783 (FLJ32783), mRNA
 NM_144969 Homo sapiens zinc finger, DHHC domain containing 15 (ZDHHC15), mRNA
 NM_144970 Homo sapiens hypothetical protein MGC39350 (MGC39350), mRNA
 NM_144972 Homo sapiens lactate dehydrogenase A-like 6A (LDHAL6A), mRNA
 NM_144973 Homo sapiens hypothetical protein MGC24039 (MGC24039), mRNA
 NM_144974 Homo sapiens hypothetical protein FLJ31846 (FLJ31846), mRNA
 NM_144975 Homo sapiens hypothetical protein MGC19764 (MGC19764), mRNA
 NM_144976 Homo sapiens zinc finger protein 564 (ZNF564), mRNA
 NM_144977 Homo sapiens family with sequence similarity 31, member B (FAM31B), mRf
 NM_144978 Homo sapiens hypothetical protein FLJ32745 (FLJ32745), mRNA
 NM_144979 Homo sapiens hypothetical protein MGC27016 (MGC27016), mRNA
 NM_144980 Homo sapiens chromosome 6 open reading frame 118 (C6orf118), mRNA
 NM_144981 Homo sapiens hypothetical protein FLJ25059 (FLJ25059), mRNA
 NM_144982 Homo sapiens hypothetical protein MGC23401 (MGC23401), mRNA
 NM_144984 Homo sapiens chromosome 10 open reading frame 72 (C10orf72), mRNA
 NM_144985 Homo sapiens cadherin-like 24 (CDH24), mRNA
 NM_144987 Homo sapiens U2(RNU2) small nuclear RNA auxiliary factor 1-like 3 (U2AF1
 NM_144988 Homo sapiens hypothetical protein MGC19780 (MGC19780), mRNA
 NM_144990 Homo sapiens hypothetical protein FLJ23878 (FLJ23878), mRNA
 NM_144991 Homo sapiens chromosome 21 open reading frame 29 (C21orf29), mRNA
 NM_144992 Homo sapiens hypothetical protein MGC26733 (MGC26733), mRNA
 NM_144994 Homo sapiens ankyrin repeat domain 23 (ANKRD23), mRNA
 NM_144995 Homo sapiens DEAH (Asp-Glu-Ala-Asp/His) box polypeptide 57 (DHX57), tr
 NM_144998 Homo sapiens hypothetical protein DKFZp761H079 (DKFZp761H079), trans
 NM_144997 Homo sapiens folliculin (BHD), transcript variant 1, mRNA
 NM_144998 Homo sapiens stimulated by retinoic acid 13 (STRA13), mRNA
 NM_144999 Homo sapiens hypothetical protein MGC20806 (MGC20806), mRNA
 NM_145000 Homo sapiens hypothetical protein FLJ25422 (FLJ25422), mRNA
 NM_145001 Homo sapiens serine/threonine kinase 32A (STK32A), mRNA
 NM_145003 Homo sapiens hypothetical protein FLJ31164 (FLJ31164), mRNA
 NM_145004 Homo sapiens a disintegrin and metalloproteinase domain 32 (ADAM32), mF
 NM_145005 Homo sapiens chromosome 9 open reading frame 72 (C9orf72), transcript v
 NM_145006 Homo sapiens sushi domain containing 3 (SUSD3), mRNA
 NM_145007 Homo sapiens NACHT, leucine rich repeat and PYD containing 11 (NALP11)
 NM_145008 Homo sapiens hypothetical protein FLJ30213 (FLJ30213), mRNA
 NM_145010 Homo sapiens chromosome 10 open reading frame 63 (C10orf63), mRNA
 NM_145011 Homo sapiens zinc finger protein 25 (KOX 19) (ZNF25), mRNA
 NM_145012 Homo sapiens chromosome 10 open reading frame 9 (C10orf9), mRNA
 NM_145013 Homo sapiens hypothetical protein MGC35558 (MGC35558), mRNA
 NM_145014 Homo sapiens hypothetical protein FLJ32915 (FLJ32915), mRNA
 NM_145015 Homo sapiens MAS-related GPR, member F (MRGPRF), mRNA
 NM_145016 Homo sapiens BXMAS2-10 (BXMAS2-10), mRNA
 NM_145017 Homo sapiens IIIg9 protein (FLJ32771), mRNA
 NM_145018 Homo sapiens hypothetical protein FLJ25416 (FLJ25416), mRNA
 NM_145019 Homo sapiens hypothetical protein FLJ30707 (FLJ30707), mRNA
 NM_145020 Homo sapiens hypothetical protein FLJ32743 (FLJ32743), mRNA
 NM_145021 Homo sapiens c-mir, cellular modulator of immune recognition (MIR), transcr
 NM_145023 Homo sapiens coiled-coil domain containing 7 (CCDC7), mRNA
 NM_145024 Homo sapiens hypothetical protein FLJ31547 (FLJ31547), mRNA
 NM_145025 Homo sapiens chromosome 6 open reading frame 199 (C6orf199), mRNA
 NM_145026 Homo sapiens spermatogenesis associated, serine-rich 1 (SPATS1), mRNA
 NM_145027 Homo sapiens chromosome 6 open reading frame 102 (C6orf102), mRNA
 NM_145028 Homo sapiens chromosome 6 open reading frame 81 (C6orf81), mRNA
 NM_145029 Homo sapiens chromosome 6 open reading frame 136 (C6orf136), mRNA
 NM_145030 Homo sapiens hypothetical protein MGC22793 (MGC22793), mRNA

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NM_145032 Homo sapiens F-box and leucine-rich repeat protein 13 (FBXL13), mRNA
NM_145033 Homo sapiens chromosome 21 open reading frame 100 (C21orf100), mRNA
NM_145034 Homo sapiens AF464140 (LOC163590), mRNA
NM_145035 Homo sapiens ADMP (ADMP), mRNA
NM_145036 Homo sapiens hypothetical protein MGC33887 (MGC33887), mRNA
NM_145037 Homo sapiens hypothetical protein MGC15606 (MGC15606), mRNA
NM_145038 Homo sapiens CG10958-like (MGC16372), mRNA
NM_145039 Homo sapiens hypothetical protein MGC16385 (MGC16385), mRNA
NM_145040 Homo sapiens protein kinase C, delta binding protein (PRKCDBP), mRNA
NM_145041 Homo sapiens hypothetical protein MGC20235 (MGC20235), mRNA
NM_145042 Homo sapiens alpha tubulin-like (MGC16703), mRNA
NM_145043 Homo sapiens nei like 2 (E. coli) (NEIL2), mRNA
NM_145044 Homo sapiens zinc finger protein 501 (ZNF501), mRNA
NM_145045 Homo sapiens hypothetical protein MGC20983 (MGC20983), mRNA
NM_145046 Homo sapiens calreticulin 3 (CALR3), mRNA
NM_145047 Homo sapiens oxidored-nitro domain-containing protein (NOR1), transcript v
NM_145048 Homo sapiens hypothetical protein MGC29898 (MGC29898), mRNA
NM_145049 Homo sapiens hypothetical protein MGC10067 (MGC10067), mRNA
NM_145050 Homo sapiens hypothetical protein MGC27434 (MGC27434), mRNA
NM_145051 Homo sapiens hypothetical protein MGC4734 (MGC4734), mRNA
NM_145052 Homo sapiens hypothetical protein MGC23937 similar to CG4798 (MGC23937), mRNA
NM_145053 Homo sapiens hypothetical protein MGC20470 (MGC20470), mRNA
NM_145054 Homo sapiens hypothetical protein LOC146845 (LOC146845), mRNA
NM_145055 Homo sapiens chromosome 18 open reading frame 25 (C18orf25), mRNA
NM_145056 Homo sapiens thymus expressed gene 3-like (MGC15476), mRNA
NM_145057 Homo sapiens CDC42 effector protein (Rho GTPase binding) 5 (CDC42EP5), mRNA
NM_145058 Homo sapiens hypothetical protein MGC7036 (MGC7036), mRNA
NM_145059 Homo sapiens fucokinase (FUK), mRNA
NM_145060 Homo sapiens chromosome 18 open reading frame 24 (C18orf24), mRNA
NM_145061 Homo sapiens chromosome 13 open reading frame 3 (C13orf3), mRNA
NM_145062 Homo sapiens chromosome 6 open reading frame 113 (C6orf113), mRNA
NM_145063 Homo sapiens chromosome 6 open reading frame 130 (C6orf130), mRNA
NM_145064 Homo sapiens SH3 and cysteine rich domain 3 (STAC3), mRNA
NM_145065 Homo sapiens pellino 3 alpha (MGC35521), mRNA
NM_145068 Homo sapiens transient receptor potential cation channel, subfamily V, mem
NM_145071 Homo sapiens cytokine inducible SH2-containing protein (CISH), transcript v
NM_145074 Homo sapiens protease, serine, 25 (PRSS25), nuclear gene encoding mitot
NM_145080 Homo sapiens non-SMC (structural maintenance of chromosomes) element
NM_145102 Homo sapiens zinc finger protein 95 homolog (mouse) (ZFP95), transcript va
NM_145109 Homo sapiens mitogen-activated protein kinase kinase 3 (MAP2K3), transcript
NM_145110 Homo sapiens mitogen-activated protein kinase kinase 3 (MAP2K3), transcript
NM_145111 Homo sapiens hypothetical protein DKFzp727G131 (DKFzp727G131), mRNA
NM_145112 Homo sapiens MAX protein (MAX), transcript variant 2, mRNA
NM_145113 Homo sapiens MAX protein (MAX), transcript variant 3, mRNA
NM_145114 Homo sapiens MAX protein (MAX), transcript variant 4, mRNA
NM_145115 Homo sapiens zinc finger protein 498 (ZNF498), mRNA
NM_145116 Homo sapiens MAX protein (MAX), transcript variant 5, mRNA
NM_145117 Homo sapiens neuron navigator 2 (NAV2), transcript variant 2, mRNA
NM_145119 Homo sapiens praja 1 (PJA1), mRNA
NM_145159 Homo sapiens jagged 2 (JAG2), transcript variant 2, mRNA
NM_145160 Homo sapiens mitogen-activated protein kinase kinase 5 (MAP2K5), transcript
NM_145161 Homo sapiens mitogen-activated protein kinase kinase 5 (MAP2K5), transcript
NM_145162 Homo sapiens mitogen-activated protein kinase kinase 5 (MAP2K5), transcript
NM_145165 Homo sapiens churchill domain containing 1 (CHURC1), mRNA
NM_145166 Homo sapiens hypothetical protein KIAA1190 (KIAA1190), mRNA
NM_145167 Homo sapiens phosphatidylinositol glycan, class M (PIGM), mRNA
NM_145168 Homo sapiens NAD(P) dependent steroid dehydrogenase-like (HSPC105), n

NM_145169 Homo sapiens chromosome 6 open reading frame 83 (C6orf83), mRNA
 NM_145170 Homo sapiens tetrapeptide repeat domain 18 (TTC18), mRNA
 NM_145171 Homo sapiens glycoprotein hormone beta 5 (GPHB5), mRNA
 NM_145172 Homo sapiens testis development protein NYD-SP29 (NYD-SP29), mRNA
 NM_145173 Homo sapiens DIRAS family, GTP-binding RAS-like 1 (DIRAS1), mRNA
 NM_145174 Homo sapiens DnaJ (Hsp40) homolog, subfamily B, member 7 (DNAJB7), m
 NM_145175 Homo sapiens NSE1 (NSE1), mRNA
 NM_145176 Homo sapiens solute carrier family 2 (facilitated glucose transporter), membe
 NM_145177 Homo sapiens dehydrogenase/reductase (SDR family) X-linked (DHRX), m
 NM_145178 Homo sapiens atonal homolog 7 (Drosophila) (ATOH7), mRNA
 NM_145179 Homo sapiens chromosome 21 open reading frame 93 (C21orf93), mRNA
 NM_145180 Homo sapiens chromosome 21 open reading frame 94 (C21orf94), mRNA
 NM_145182 Homo sapiens PYD and CARD domain containing (PYCARD), transcript vari
 NM_145183 Homo sapiens PYD and CARD domain containing (PYCARD), transcript vari
 NM_145185 Homo sapiens mitogen-activated protein kinase kinase 7 (MAP2K7), mRNA
 NM_145186 Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 11
 NM_145187 Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 12
 NM_145188 Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 12
 NM_145189 Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 12
 NM_145190 Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 12
 NM_145196 Homo sapiens lipoyltransferase 1 (LIPT1), transcript variant 2, mRNA
 NM_145197 Homo sapiens lipoyltransferase 1 (LIPT1), transcript variant 3, mRNA
 NM_145198 Homo sapiens lipoyltransferase 1 (LIPT1), transcript variant 4, mRNA
 NM_145199 Homo sapiens lipoyltransferase 1 (LIPT1), transcript variant 5, mRNA
 NM_145200 Homo sapiens calcium binding protein 4 (CABP4), mRNA
 NM_145201 Homo sapiens similar to CG3714 gene product (PP3858), mRNA
 NM_145202 Homo sapiens proline-rich acidic protein 1 (PRAP1), mRNA
 NM_145203 Homo sapiens casein kinase 1, alpha 1-like (CSNK1A1L), mRNA
 NM_145204 Homo sapiens SUMO/sentrin specific protease family member 8 (SENP8), m
 NM_145205 Homo sapiens HMG2 like (LOC127540), mRNA
 NM_145206 Homo sapiens vesicle transport through interaction with t-SNAREs homolog
 NM_145207 Homo sapiens spermatogenesis associated 5 (SPATA5), mRNA
 NM_145208 Homo sapiens methyl-CpG binding domain protein 3-like 1 (MBD3L1), mRN
 NM_145212 Homo sapiens mitochondrial ribosomal protein L30 (MRPL30), nuclear gene
 NM_145213 Homo sapiens mitochondrial ribosomal protein L30 (MRPL30), nuclear gene
 NM_145214 Homo sapiens tripartite motif-containing 11 (TRIM11), mRNA
 NM_145230 Homo sapiens chromosome 7 open reading frame 32 (C7orf32), mRNA
 NM_145231 Homo sapiens chromosome 14 open reading frame 143 (C14orf143), mRNA
 NM_145232 Homo sapiens LOC90353 (LOC90353), mRNA
 NM_145233 Homo sapiens zinc finger protein 625 (ZNF625), mRNA
 NM_145234 Homo sapiens chordin-like 1 (CHRD1), mRNA
 NM_145235 Homo sapiens fibronectin type 3 and ankyrin repeat domains 1 (FANK1), mR
 NM_145236 Homo sapiens UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransfera
 NM_145237 Homo sapiens similar to RNA polymerase I transcription factor RRN3 (LOC9
 NM_145238 Homo sapiens zinc finger protein 31 (KOX 29) (ZNF31), mRNA
 NM_145239 Homo sapiens similar to lymphocyte antigen 6 complex, locus G5B; G5b pro
 NM_145241 Homo sapiens WD repeat domain 31 (WDR31), mRNA
 NM_145242 Homo sapiens similar to POSSIBLE GUSTATORY RECEPTOR CLONE PTE
 NM_145243 Homo sapiens metalloprotease related protein 1 (MPRP-1), mRNA
 NM_145244 Homo sapiens DNA-damage-inducible transcript 4-like (DDIT4L), mRNA
 NM_145245 Homo sapiens ecotropic viral integration site 5-like (EVL), mRNA
 NM_145246 Homo sapiens chromosome 10 open reading frame 4 (C10orf4), transcript ve
 NM_145247 Homo sapiens chromosome 10 open reading frame 78 (C10orf78), transcript
 NM_145248 Homo sapiens LOC122258 (LOC122258), mRNA
 NM_145249 Homo sapiens family with sequence similarity 14, member B (FAM14B), mR
 NM_145250 Homo sapiens chromosome 14 open reading frame 6 (C14orf6), mRNA
 NM_145251 Homo sapiens serine/threonine/tyrosine interacting protein (STYX), mRNA

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NM_145252 Homo sapiens similar to common salivary protein 1 (LOC124220), mRNA
NM_145253 Homo sapiens LOC124402 (LOC124402), mRNA
NM_145254 Homo sapiens LOC124491 (LOC124491), mRNA
NM_145255 Homo sapiens mitochondrial ribosomal protein L10 (MRPL10), nuclear gene
NM_145256 Homo sapiens leucine rich repeat containing 25 (LRRC25), mRNA
NM_145257 Homo sapiens LOC126731 (LOC126731), mRNA
NM_145258 Homo sapiens hypothetical protein MGC22773 (MGC22773), mRNA
NM_145259 Homo sapiens activin A receptor, type IC (ACVR1C), mRNA
NM_145260 Homo sapiens odd-skipped homolog (Drosophila) (ODD), mRNA
NM_145261 Homo sapiens homolog of yeast TIM14 (TIM14), transcript variant 1, mRNA
NM_145262 Homo sapiens CG9886-like (GLYCTK), mRNA
NM_145263 Homo sapiens LOC132671 (LOC132671), mRNA
NM_145265 Homo sapiens similar to RIKEN cDNA 0610011N22 (LOC133957), mRNA
NM_145266 Homo sapiens similar to RIKEN cDNA 2700047N05 (LOC134492), mRNA
NM_145267 Homo sapiens chromosome 6 open reading frame 57 (C6orf57), mRNA
NM_145268 Homo sapiens LOC136263 (LOC136263), mRNA
NM_145269 Homo sapiens similar to CG6405 gene product (LOC137392), mRNA
NM_145270 Homo sapiens similar to hypothetical protein FLJ13841 (LOC146325), mRNA
NM_145271 Homo sapiens similar to hypothetical protein MGC13138 (LOC146542), mRNA
NM_145272 Homo sapiens LOC146853 (LOC146853), mRNA
NM_145273 Homo sapiens triggering receptor expressed on myeloid cells 4 (TREM4), mRNA
NM_145274 Homo sapiens hypothetical protein MGC21518 (MGC21518), mRNA
NM_145275 Homo sapiens kinesin light chain 2-like (KLC2L), transcript variant 2, mRNA
NM_145276 Homo sapiens zinc finger protein 563 (ZNF563), mRNA
NM_145277 Homo sapiens hemochromatosis type 2 (juvenile) (HFE2), transcript variant 1
NM_145278 Homo sapiens LOC148823 (LOC148823), mRNA
NM_145279 Homo sapiens MOB1, Mps One Binder kinase activator-like 2C (yeast) (MOE)
NM_145280 Homo sapiens similar to hepatocellular carcinoma-associated antigen HCA5
NM_145282 Homo sapiens similar to CG4995 gene product (LOC153328), mRNA
NM_145283 Homo sapiens chromosome 9 open reading frame 121 (C9orf121), mRNA
NM_145284 Homo sapiens similar to hypothetical protein MGC17347 (LOC159090), mRNA
NM_145285 Homo sapiens NK2 transcription factor related, locus 3 (Drosophila) (NKX2-3)
NM_145286 Homo sapiens stomatin (EPB72)-like 3 (STOML3), mRNA
NM_145287 Homo sapiens zinc finger protein 519 (ZNF519), mRNA
NM_145288 Homo sapiens zinc finger protein 342 (ZNF342), mRNA
NM_145291 Homo sapiens zinc finger protein 509 (ZNF509), mRNA
NM_145292 Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgl
NM_145293 Homo sapiens similar to hypothetical protein FLJ20897 (LOC196549), mRNA
NM_145294 Homo sapiens similar to RIKEN cDNA 3230401M21 [Mus musculus] (LOC19
NM_145295 Homo sapiens zinc finger protein 627 (ZNF627), mRNA
NM_145296 Homo sapiens immunoglobulin superfamily, member 4C (IGSF4C), mRNA
NM_145297 Homo sapiens zinc finger protein 626 (ZNF626), mRNA
NM_145298 Homo sapiens apolipoprotein B mRNA editing enzyme, catalytic polypeptide
NM_145299 Homo sapiens similar to Dynein heavy chain at 16F (LOC200383), mRNA
NM_145300 Homo sapiens LOC200420 (LOC200420), mRNA
NM_145301 Homo sapiens similar to CG-148 protein (LOC201158), mRNA
NM_145303 Homo sapiens similar to RIKEN cDNA 2310008M10 (LOC202459), mRNA
NM_145304 Homo sapiens chromosome 7 open reading frame 33 (C7orf33), mRNA
NM_145305 Homo sapiens similar to solute carrier family 25, member 16 (LOC203427),
NM_145306 Homo sapiens chromosome 10 open reading frame 35 (C10orf35), mRNA
NM_145307 Homo sapiens pleckstrin homology domain containing, family K member 1 (F
NM_145308 Homo sapiens hypothetical protein BC004224 (LOC220070), mRNA
NM_145309 Homo sapiens Hypothetical 55.1 kDa protein F09G8.5 in chromosome III (LC
NM_145310 Homo sapiens zinc finger protein 258 (ZNF258), mRNA
NM_145311 Homo sapiens crystallin, zeta (quinone reductase)-like 1 (CRYZL1), transcript
NM_145312 Homo sapiens zinc finger protein 485 (ZNF485), mRNA
NM_145313 Homo sapiens RasGEF domain family, member 1A (RASGEF1A), mRNA

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NM_145314 Homo sapiens chromosome 10 open reading frame 49 (C10orf49), mRNA
 NM_145315 Homo sapiens lactation elevated 1 (LACE1), mRNA
 NM_145316 Homo sapiens chromosome 6 open reading frame 128 (C6orf128), mRNA
 NM_145320 Homo sapiens oxysterol binding protein-like 3 (OSBPL3), transcript variant 2
 NM_145321 Homo sapiens oxysterol binding protein-like 3 (OSBPL3), transcript variant 3
 NM_145322 Homo sapiens oxysterol binding protein-like 3 (OSBPL3), transcript variant 4
 NM_145323 Homo sapiens oxysterol binding protein-like 3 (OSBPL3), transcript variant 5
 NM_145324 Homo sapiens oxysterol binding protein-like 3 (OSBPL3), transcript variant 6
 NM_145325 Homo sapiens DNA directed RNA polymerase II polypeptide J-related gene 1
 NM_145326 Homo sapiens similar to hypothetical protein FLJ13659 (LOC115648), mRNA
 NM_145328 Homo sapiens chromosome 21 open reading frame 66 (C21orf66), transcript
 NM_145330 Homo sapiens mitochondrial ribosomal protein L33 (MRPL33), nuclear gene
 NM_145331 Homo sapiens mitogen-activated protein kinase kinase kinase 7 (MAP3K7), l
 NM_145332 Homo sapiens mitogen-activated protein kinase kinase kinase 7 (MAP3K7), l
 NM_145333 Homo sapiens mitogen-activated protein kinase kinase kinase 7 (MAP3K7), l
 NM_145341 Homo sapiens programmed cell death 4 (neoplastic transformation inhibitor)
 NM_145342 Homo sapiens mitogen-activated protein kinase kinase kinase 7 interacting f
 NM_145343 Homo sapiens apolipoprotein L, 1 (APOL1), transcript variant 2, mRNA
 NM_145344 Homo sapiens apolipoprotein L, 1 (APOL1), transcript variant 3, mRNA
 NM_145345 Homo sapiens socius (SOC), mRNA
 NM_145346 Homo sapiens socius (SOC), mRNA
 NM_145347 Homo sapiens kringle containing transmembrane protein 2 (KREMEN2), tran
 NM_145348 Homo sapiens kringle containing transmembrane protein 2 (KREMEN2), tran
 NM_145349 Homo sapiens scavenger receptor class F, member 1 (SCARF1), transcript v
 NM_145350 Homo sapiens scavenger receptor class F, member 1 (SCARF1), transcript v
 NM_145351 Homo sapiens scavenger receptor class F, member 1 (SCARF1), transcript v
 NM_145352 Homo sapiens scavenger receptor class F, member 1 (SCARF1), transcript v
 NM_145637 Homo sapiens apolipoprotein L, 2 (APOL2), transcript variant beta, mRNA
 NM_145638 Homo sapiens oxysterol binding protein-like 5 (OSBPL5), transcript variant 2
 NM_145639 Homo sapiens apolipoprotein L, 3 (APOL3), transcript variant alpha/c, mRNA
 NM_145640 Homo sapiens apolipoprotein L, 3 (APOL3), transcript variant alpha/d, mRNA
 NM_145641 Homo sapiens apolipoprotein L, 3 (APOL3), transcript variant beta/a, mRNA
 NM_145642 Homo sapiens apolipoprotein L, 3 (APOL3), transcript variant beta/b, mRNA
 NM_145644 Homo sapiens mitochondrial ribosomal protein L35 (MRPL35), nuclear gene
 NM_145645 Homo sapiens Williams-Beuren Syndrome critical region protein 20 copy B (C
 NM_145646 Homo sapiens DEAH (Asp-Glu-Ala-Asp/His) box polypeptide 57 (DHX57), tr
 NM_145647 Homo sapiens unknown MGC21854 product (MGC21854), mRNA
 NM_145648 Homo sapiens solute carrier family 15, member 4 (SLC15A4), mRNA
 NM_145649 Homo sapiens glucosaminyl (N-acetyl) transferase 2, I-branching enzyme (G
 NM_145650 Homo sapiens mucin 15 (MUC15), mRNA
 NM_145651 Homo sapiens ligand binding protein RYD5 (RYD5), mRNA
 NM_145652 Homo sapiens WAP four-disulfide core domain 5 (WFDC5), mRNA
 NM_145653 Homo sapiens transcription elongation factor B polypeptide 3C (elongin A3) (l
 NM_145654 Homo sapiens RAD52 homolog B (S. cerevisiae) (RAD52B), mRNA
 NM_145655 Homo sapiens glucosaminyl (N-acetyl) transferase 2, I-branching enzyme (G
 NM_145657 Homo sapiens GS homeobox 1 (GSH1), mRNA
 NM_145658 Homo sapiens sperm equatorial segment protein 1 (SPESP1), mRNA
 NM_145659 Homo sapiens Interleukin 27 (IL27), mRNA
 NM_145660 Homo sapiens apolipoprotein L, 4 (APOL4), transcript variant b, mRNA
 NM_145662 Homo sapiens SPANX family, member A2 (SPANXA2), mRNA
 NM_145663 Homo sapiens Dbf4-related factor 1 (DRF1), transcript variant 1, mRNA
 NM_145664 Homo sapiens SPANX family, member B2 (SPANXB2), mRNA
 NM_145665 Homo sapiens SPANX family, member E (SPANXE), mRNA
 NM_145685 Homo sapiens BRP1 homolog, subunit of RNA polymerase III transcription in
 NM_145686 Homo sapiens mitogen-activated protein kinase kinase kinase kinase 4 (MAF
 NM_145687 Homo sapiens mitogen-activated protein kinase kinase kinase kinase 4 (MAF
 NM_145689 Homo sapiens amyloid beta (A4) precursor protein-binding, family B, membe

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NM_145690 Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase acti
 NM_145691 Homo sapiens ATP synthase mitochondrial F1 complex assembly factor 2 (A
 NM_145693 Homo sapiens lipin 1 (LPIN1), mRNA
 NM_145695 Homo sapiens diacylglycerol kinase, beta 90kDa (DGKB), transcript variant 2
 NM_145696 Homo sapiens BRF1 homolog, subunit of RNA polymerase III transcription in
 NM_145697 Homo sapiens cell division cycle associated 1 (CDCA1), transcript variant 1,
 NM_145698 Homo sapiens acyl-Coenzyme A binding domain containing 5 (ACBD5), mR
 NM_145699 Homo sapiens apolipoprotein B mRNA editing enzyme, catalytic polypeptide
 NM_145701 Homo sapiens cell division cycle associated 4 (CDCA4), transcript variant 2,
 NM_145702 Homo sapiens tigger transposable element derived 1 (TIGD1), mRNA
 NM_145714 Homo sapiens ataxin 2 related protein (A2LP), transcript variant B, mRNA
 NM_145715 Homo sapiens tigger transposable element derived 2 (TIGD2), mRNA
 NM_145716 Homo sapiens single stranded DNA binding protein 3 (SSBP3), mRNA
 NM_145719 Homo sapiens tigger transposable element derived 3 (TIGD3), mRNA
 NM_145720 Homo sapiens tigger transposable element derived 4 (TIGD4), mRNA
 NM_145725 Homo sapiens TNF receptor-associated factor 3 (TRAF3), transcript variant
 NM_145726 Homo sapiens TNF receptor-associated factor 3 (TRAF3), transcript variant
 NM_145727 Homo sapiens lipoprotein, Lp(a)-like 2 (LPAL2), transcript variant 2, mRNA
 NM_145728 Homo sapiens desmuslin (DMN), transcript variant A, mRNA
 NM_145729 Homo sapiens mitochondrial ribosomal protein L24 (MRPL24), nuclear gene
 NM_145730 Homo sapiens adaptor-related protein complex 1, beta 1 subunit (AP1B1), tr
 NM_145731 Homo sapiens synaptogyrin 1 (SYNGR1), transcript variant 1b, mRNA
 NM_145733 Homo sapiens septin 3 (SEPT3), transcript variant A, mRNA
 NM_145734 Homo sapiens septin 3 (SEPT3), transcript variant C, mRNA
 NM_145735 Homo sapiens Rho guanine nucleotide exchange factor (GEF) 7 (ARHGEF7,
 NM_145738 Homo sapiens synaptogyrin 1 (SYNGR1), transcript variant 1c, mRNA
 NM_145739 Homo sapiens oxysterol binding protein-like 6 (OSBPL6), transcript variant 2
 NM_145740 Homo sapiens glutathione S-transferase A1 (GSTA1), mRNA
 NM_145747 Homo sapiens thioredoxin reductase 2 (TXNRD2), nuclear gene encoding m
 NM_145748 Homo sapiens thioredoxin reductase 2 (TXNRD2), nuclear gene encoding m
 NM_145751 Homo sapiens TNF receptor-associated factor 4 (TRAF4), transcript variant
 NM_145752 Homo sapiens CDP-diacylglycerol--inositol 3-phosphatidyltransferase (phos
 NM_145753 Homo sapiens plectstrin homology-like domain, family B, member 2 (PHLDE
 NM_145764 Homo sapiens kinesin family member C2 (KIFC2), mRNA
 NM_145755 Homo sapiens TPR domain containing STI2 (STI2), mRNA
 NM_145756 Homo sapiens zinc finger protein 396 (ZNF396), mRNA
 NM_145759 Homo sapiens TNF receptor-associated factor 5 (TRAF5), transcript variant
 NM_145762 Homo sapiens GDNF family receptor alpha 4 (GFRA4), transcript variant 2, r
 NM_145763 Homo sapiens GDNF family receptor alpha 4 (GFRA4), transcript variant 3, r
 NM_145764 Homo sapiens microsomal glutathione S-transferase 1 (MGST1), transcript v
 NM_145791 Homo sapiens microsomal glutathione S-transferase 1 (MGST1), transcript v
 NM_145792 Homo sapiens microsomal glutathione S-transferase 1 (MGST1), transcript v
 NM_145793 Homo sapiens GDNF family receptor alpha 1 (GFRA1), transcript variant 2, r
 NM_145794 Homo sapiens downstream neighbor of SON (DONSON), transcript variant 2
 NM_145795 Homo sapiens downstream neighbor of SON (DONSON), transcript variant 3
 NM_145796 Homo sapiens pogo transposable element with ZNF domain (POGZ), transc
 NM_145798 Homo sapiens oxysterol binding protein-like 7 (OSBPL7), transcript variant 1
 NM_145799 Homo sapiens septin 6 (SEPT6), transcript variant 1, mRNA
 NM_145800 Homo sapiens septin 6 (SEPT6), transcript variant III, mRNA
 NM_145802 Homo sapiens septin 6 (SEPT6), transcript variant V, mRNA
 NM_145803 Homo sapiens TNF receptor-associated factor 6 (TRAF6), transcript variant
 NM_145804 Homo sapiens ankyrin repeat and BTB (POZ) domain containing 2 (ABTB2),
 NM_145805 Homo sapiens ISL2 transcription factor, LIM/homeodomain, (Islet-2) (ISL2), r
 NM_145806 Homo sapiens zinc finger protein 511 (ZNF511), mRNA
 NM_145807 Homo sapiens hypothetical protein BC018697 (LOC126147), mRNA
 NM_145808 Homo sapiens myotrophin (MTPN), mRNA
 NM_145809 Homo sapiens TL132 protein (LOC220594), mRNA

NM_145810 Homo sapiens cell division cycle associated 7 (CDCA7), transcript variant 2,
 NM_145811 Homo sapiens calcium channel, voltage-dependent, gamma subunit 5 (CACI
 NM_145812 Homo sapiens programmed cell death 8 (apoptosis-inducing factor) (PDCD8
 NM_145813 Homo sapiens programmed cell death 8 (apoptosis-inducing factor) (PDCD8
 NM_145814 Homo sapiens calcium channel, voltage-dependent, gamma subunit 6 (CACI
 NM_145815 Homo sapiens calcium channel, voltage-dependent, gamma subunit 6 (CACI
 NM_145818 Homo sapiens component of oligomeric golgi complex 4 (COG4), transcript v
 NM_145858 Homo sapiens crystallin, zeta (quinone reductase)-like 1 (CRYZL1), transcrip
 NM_145859 Homo sapiens programmed cell death 10 (PDCD10), transcript variant 2, mF
 NM_145860 Homo sapiens programmed cell death 10 (PDCD10), transcript variant 3, mF
 NM_145861 Homo sapiens EDAR-associated death domain (EDARADD), transcript varia
 NM_145862 Homo sapiens CHK2 checkpoint homolog (S. pombe) (CHEK2), transcript va
 NM_145863 Homo sapiens ankyrin repeat and SOCS box-containing 3 (ASB3), transcript
 NM_145864 Homo sapiens kallikrein 3, (prostate specific antigen) (KLK3), transcript varia
 NM_145865 Homo sapiens hypothetical protein FLJ38819 (FLJ38819), mRNA
 NM_145867 Homo sapiens leukotriene C4 synthase (LTC4S), transcript variant 1, mRNA
 NM_145868 Homo sapiens annexin A11 (ANXA11), transcript variant b, mRNA
 NM_145869 Homo sapiens annexin A11 (ANXA11), transcript variant c, mRNA
 NM_145870 Homo sapiens glutathione transferase zeta 1 (maleylacetoacetate isomerase
 NM_145871 Homo sapiens glutathione transferase zeta 1 (maleylacetoacetate isomerase
 NM_145872 Homo sapiens ankyrin repeat and SOCS box-containing 4 (ASB4), transcript
 NM_145886 Homo sapiens leucine-rich repeats and death domain containing (LRDD), tra
 NM_145887 Homo sapiens leucine-rich repeats and death domain containing (LRDD), tra
 NM_145888 Homo sapiens kallikrein 10 (KLK10), transcript variant 2, mRNA
 NM_145891 Homo sapiens ataxin 2-binding protein 1 (A2BP1), transcript variant 1, mRN/
 NM_145892 Homo sapiens ataxin 2-binding protein 1 (A2BP1), transcript variant 2, mRN/
 NM_145893 Homo sapiens ataxin 2-binding protein 1 (A2BP1), transcript variant 3, mRN/
 NM_145894 Homo sapiens kallikrein 12 (KLK12), transcript variant 2, mRNA
 NM_145895 Homo sapiens kallikrein 12 (KLK12), transcript variant 3, mRNA
 NM_145896 Homo sapiens prefoldin 5 (PFDN5), transcript variant 2, mRNA
 NM_145897 Homo sapiens prefoldin 5 (PFDN5), transcript variant 3, mRNA
 NM_145898 Homo sapiens chemokine (C-C motif) ligand 23 (CCL23), transcript variant C
 NM_145899 Homo sapiens high mobility group AT-hook 1 (HMGA1), transcript variant 1, i
 NM_145901 Homo sapiens high mobility group AT-hook 1 (HMGA1), transcript variant 3, i
 NM_145902 Homo sapiens high mobility group AT-hook 1 (HMGA1), transcript variant 4, i
 NM_145903 Homo sapiens high mobility group AT-hook 1 (HMGA1), transcript variant 5, i
 NM_145904 Homo sapiens high mobility group AT-hook 1 (HMGA1), transcript variant 6, i
 NM_145905 Homo sapiens high mobility group AT-hook 1 (HMGA1), transcript variant 7, i
 NM_145906 Homo sapiens RIO kinase 3 (yeast) (RIOK3), transcript variant 2, mRNA
 NM_145909 Homo sapiens zinc finger protein 323 (ZNF323), mRNA
 NM_145910 Homo sapiens NIMA (never in mitosis gene a)-related kinase 11 (NEK11), m
 NM_145911 Homo sapiens zinc finger protein 23 (KOX 16) (ZNF23), mRNA
 NM_145912 Homo sapiens NFAT activation molecule 1 (NFAM1), mRNA
 NM_145913 Homo sapiens solute carrier family 5 (iodide transporter), member 8 (SLC5A/
 NM_145914 Homo sapiens zinc finger protein 38 (ZNF38), mRNA
 NM_145918 Homo sapiens cathepsin L (CTSL), transcript variant 2, mRNA
 NM_146387 Homo sapiens mitochondrial ribosomal protein L4 (MRPL4), nuclear gene en
 NM_146388 Homo sapiens mitochondrial ribosomal protein L4 (MRPL4), nuclear gene en
 NM_146421 Homo sapiens glutathione S-transferase M1 (GSTM1), transcript variant 2, r
 NM_147127 Homo sapiens Ellis van Creveld syndrome 2 (limbin) (EVC2), mRNA
 NM_147128 Homo sapiens zinc and ring finger 2 (ZNRF2), mRNA
 NM_147129 Homo sapiens hypothetical protein LOC259173 (FLJ36525), transcript variat
 NM_147130 Homo sapiens natural cytotoxicity triggering receptor 3 (NCR3), mRNA
 NM_147131 Homo sapiens galactose-1-phosphate uridylyltransferase (GALT), transcript
 NM_147132 Homo sapiens galactose-1-phosphate uridylyltransferase (GALT), transcript
 NM_147133 Homo sapiens nuclear transcription factor, X-box binding 1 (NFX1), transcrip
 NM_147134 Homo sapiens nuclear transcription factor, X-box binding 1 (NFX1), transcrip

NM_147147 Homo sapiens blood vessel epicardial substance (BVES), transcript variant E
 NM_147148 Homo sapiens glutathione S-transferase M4 (GSTM4), transcript variant 2, r
 NM_147149 Homo sapiens glutathione S-transferase M4 (GSTM4), transcript variant 3, r
 NM_147150 Homo sapiens A kinase (PRKA) anchor protein 2 (AKAP2), transcript variant
 NM_147152 Homo sapiens intersectin 2 (ITSN2), transcript variant 2, mRNA
 NM_147156 Homo sapiens transmembrane protein 23 (TMEM23), mRNA
 NM_147157 Homo sapiens opioid receptor, sigma 1 (OPRS1), transcript variant 2, mRNA
 NM_147158 Homo sapiens opioid receptor, sigma 1 (OPRS1), transcript variant 3, mRNA
 NM_147159 Homo sapiens opioid receptor, sigma 1 (OPRS1), transcript variant 4, mRNA
 NM_147160 Homo sapiens opioid receptor, sigma 1 (OPRS1), transcript variant 5, mRNA
 NM_147161 Homo sapiens thioesterase, adipose associated (THEA), transcript variant 2,
 NM_147162 Homo sapiens interleukin 11 receptor, alpha (IL11RA), transcript variant 2, r
 NM_147164 Homo sapiens ciliary neurotrophic factor receptor (CNTFR), transcript varian
 NM_147166 Homo sapiens A kinase (PRKA) anchor protein (yotiao) 9 (AKAP9), transcrip
 NM_147168 Homo sapiens chromosome 9 open reading frame 24 (C9orf24), transcript va
 NM_147169 Homo sapiens chromosome 9 open reading frame 24 (C9orf24), transcript va
 NM_147171 Homo sapiens A kinase (PRKA) anchor protein (yotiao) 9 (AKAP9), transcrip
 NM_147172 Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 2 (l
 NM_147173 Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 2 (l
 NM_147174 Homo sapiens heparan sulfate 6-O-sulfotransferase 2 (HS6ST2), mRNA
 NM_147175 Homo sapiens heparan sulfate 6-O-sulfotransferase 2 (HS6ST2), transcript v
 NM_147180 Homo sapiens protein phosphatase 3 (formerly 2B), regulatory subunit B, 19l
 NM_147181 Homo sapiens Kv channel interacting protein 4 (KCNP4), transcript variant 2
 NM_147182 Homo sapiens Kv channel interacting protein 4 (KCNP4), transcript variant 3
 NM_147183 Homo sapiens Kv channel interacting protein 4 (KCNP4), transcript variant 4
 NM_147184 Homo sapiens tumor protein p53 inducible protein 3 (TP53I3), transcript vari
 NM_147185 Homo sapiens A kinase (PRKA) anchor protein (yotiao) 9 (AKAP9), transcrip
 NM_147187 Homo sapiens tumor necrosis factor receptor superfamily, member 10b (TNF
 NM_147188 Homo sapiens F-box protein 22 (FBXO22), transcript variant 1, mRNA
 NM_147189 Homo sapiens hypothetical protein MGC39325 (MGC39325), mRNA
 NM_147190 Homo sapiens LAG1 longevity assurance homolog 5 (S. cerevisiae) (LASS5)
 NM_147191 Homo sapiens matrix metalloproteinase 21 (MMP21), mRNA
 NM_147192 Homo sapiens diencephalon/mesencephalon homeobox 1 (DMBX1), transcri
 NM_147193 Homo sapiens GLIS family zinc finger 1 (GLIS1), mRNA
 NM_147194 Homo sapiens hypothetical protein MGC35361 (MGC35361), mRNA
 NM_147195 Homo sapiens FLJ35740 protein (FLJ35740), mRNA
 NM_147196 Homo sapiens transmembrane inner ear (TMIE), mRNA
 NM_147197 Homo sapiens WAP four-disulfide core domain 11 (WFDC11), mRNA
 NM_147198 Homo sapiens WAP four-disulfide core domain 9 (WFDC9), mRNA
 NM_147199 Homo sapiens G protein-coupled receptor MRGX1 (MRGX1), mRNA
 NM_147200 Homo sapiens chromosome 6 open reading frame 4 (C6orf4), transcript vari
 NM_147202 Homo sapiens chromosome 9 open reading frame 25 (C9orf25), mRNA
 NM_147203 Homo sapiens fibrinogen-like 1 (FGL1), transcript variant 2, mRNA
 NM_147204 Homo sapiens transient receptor potential cation channel, subfamily V, mem
 NM_147223 Homo sapiens nuclear receptor coactivator 1 (NCOA1), transcript variant 2, r
 NM_147233 Homo sapiens nuclear receptor coactivator 1 (NCOA1), transcript variant 3, r
 NM_147696 Homo sapiens chromosome 6 open reading frame 4 (C6orf4), transcript vari
 NM_147777 Homo sapiens sorting nexin 15 (SNX15), transcript variant B, mRNA
 NM_147780 Homo sapiens cathepsin B (CTSB), transcript variant 2, mRNA
 NM_147781 Homo sapiens cathepsin B (CTSB), transcript variant 3, mRNA
 NM_147782 Homo sapiens cathepsin B (CTSB), transcript variant 4, mRNA
 NM_147783 Homo sapiens cathepsin B (CTSB), transcript variant 5, mRNA
 NM_148169 Homo sapiens F-box protein 17 (FBXO17), transcript variant 1, mRNA
 NM_148170 Homo sapiens cathepsin C (CTSC), transcript variant 2, mRNA
 NM_148171 Homo sapiens ubiquitin associated protein 2 (UBAP2), transcript variant 3, r
 NM_148172 Homo sapiens phosphatidylethanolamine N-methyltransferase (PEMT), nucl
 NM_148173 Homo sapiens phosphatidylethanolamine N-methyltransferase (PEMT), nucl

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NM_148174 Homo sapiens ornithine decarboxylase antizyme inhibitor (OAZIN), transcript
 NM_148175 Homo sapiens peptidylprolyl isomerase (cyclophilin)-like 2 (PPIL2), transcript
 NM_148176 Homo sapiens peptidylprolyl isomerase (cyclophilin)-like 2 (PPIL2), transcript
 NM_148177 Homo sapiens F-box protein 32 (FBXO32), transcript variant 2, mRNA
 NM_148178 Homo sapiens chromosome 9 open reading frame 23 (C9orf23), transcript v2
 NM_148179 Homo sapiens chromosome 9 open reading frame 23 (C9orf23), transcript v2
 NM_148414 Homo sapiens ataxin 2 related protein (A2LP), transcript variant C, mRNA
 NM_148415 Homo sapiens ataxin 2 related protein (A2LP), transcript variant D, mRNA
 NM_148416 Homo sapiens ataxin 2 related protein (A2LP), transcript variant E, mRNA
 NM_148570 Homo sapiens mitochondrial ribosomal protein L27 (MRPL27), nuclear gene
 NM_148571 Homo sapiens mitochondrial ribosomal protein L27 (MRPL27), nuclear gene
 NM_148672 Homo sapiens chemokine (C-C motif) ligand 28 (CCL28), transcript variant 2
 NM_148674 Homo sapiens SMC1 structural maintenance of chromosomes 1-like 2 (yeast
 NM_148675 Homo sapiens Down syndrome critical region gene 9 (DSCR9), mRNA
 NM_148676 Homo sapiens Down syndrome critical region gene 10 (DSCR10), mRNA
 NM_148682 Homo sapiens Williams-Beuren syndrome chromosome region 16 (WBSCR1
 NM_148686 Homo sapiens Smith-Magenis syndrome chromosome region, candidate 7 (S
 NM_148687 Homo sapiens mitochondrial ribosomal protein L10 (MRPL10), nuclear gene
 NM_148688 Homo sapiens chemokine (C-C motif) ligand 25 (CCL25), transcript variant 2
 NM_148694 Homo sapiens family with sequence similarity 44, member A (FAM44A), mR
 NM_148696 Homo sapiens preproneurotrophin B (NPB), mRNA
 NM_148697 Homo sapiens orphan short-chain dehydrogenase / reductase (SDR-O), mR
 NM_148698 Homo sapiens forkhead box P2 (FOXP2), transcript variant 2, mRNA
 NM_148699 Homo sapiens forkhead box P2 (FOXP2), transcript variant 3, mRNA
 NM_148900 Homo sapiens forkhead box P2 (FOXP2), transcript variant 4, mRNA
 NM_148901 Homo sapiens tumor necrosis factor receptor superfamily, member 18 (TNFF
 NM_148902 Homo sapiens tumor necrosis factor receptor superfamily, member 18 (TNFF
 NM_148903 Homo sapiens GREB1 protein (GREB1), transcript variant c, mRNA
 NM_148904 Homo sapiens oxysterol binding protein-like 9 (OSBPL9), transcript variant 1
 NM_148905 Homo sapiens oxysterol binding protein-like 9 (OSBPL9), transcript variant 2
 NM_148906 Homo sapiens oxysterol binding protein-like 9 (OSBPL9), transcript variant 3
 NM_148907 Homo sapiens oxysterol binding protein-like 9 (OSBPL9), transcript variant 4
 NM_148908 Homo sapiens oxysterol binding protein-like 9 (OSBPL9), transcript variant 5
 NM_148909 Homo sapiens oxysterol binding protein-like 9 (OSBPL9), transcript variant 7
 NM_148910 Homo sapiens toll-interleukin 1 receptor (TIR) domain containing adaptor pro
 NM_148911 Homo sapiens CHRNA7 (cholinergic receptor, nicotinic, alpha polypeptide 7,
 NM_148912 Homo sapiens Williams Beuren syndrome chromosome region 21 (WBSCR2
 NM_148913 Homo sapiens Williams Beuren syndrome chromosome region 21 (WBSCR2
 NM_148914 Homo sapiens Williams Beuren syndrome chromosome region 21 (WBSCR2
 NM_148915 Homo sapiens Williams Beuren syndrome chromosome region 21 (WBSCR2
 NM_148916 Homo sapiens Williams Beuren syndrome chromosome region 21 (WBSCR2
 NM_148918 Homo sapiens serine hydroxymethyltransferase 1 (soluble) (SHMT1), transc
 NM_148919 Homo sapiens proteasome (prosome, macropain) subunit, beta type, 8 (large
 NM_148920 Homo sapiens phosphatidylinositol glycan, class Q (PIGQ), transcript variant
 NM_148921 Homo sapiens epsin 2 (EPN2), transcript variant 1, mRNA
 NM_148923 Homo sapiens cytochrome b-5 (CYB5), mRNA
 NM_148936 Homo sapiens Williams Beuren syndrome chromosome region 20C (WBSCR
 NM_148954 Homo sapiens proteasome (prosome, macropain) subunit, beta type, 9 (large
 NM_148955 Homo sapiens sorting nexin 1 (SNX1), transcript variant 2, mRNA
 NM_148956 Homo sapiens Williams Beuren syndrome chromosome region 20A (WBSCR
 NM_148957 Homo sapiens tumor necrosis factor receptor superfamily, member 19 (TNFF
 NM_148959 Homo sapiens HUS1 checkpoint homolog b (S. pombe) (HUS1B), mRNA
 NM_148960 Homo sapiens claudin 19 (CLDN19), mRNA
 NM_148961 Homo sapiens otospiralin (OTOS), mRNA
 NM_148962 Homo sapiens oxoecicosanoid (OXE) receptor 1 (OXER1), mRNA
 NM_148963 Homo sapiens G protein-coupled receptor, family C, group 6, member A (GP
 NM_148964 Homo sapiens cathepsin E (CTSE), transcript variant 2, mRNA

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NM_148965 Homo sapiens tumor necrosis factor receptor superfamily, member 25 (TNFF
 NM_148966 Homo sapiens tumor necrosis factor receptor superfamily, member 25 (TNFF
 NM_148967 Homo sapiens tumor necrosis factor receptor superfamily, member 25 (TNFF
 NM_148968 Homo sapiens tumor necrosis factor receptor superfamily, member 25 (TNFF
 NM_148969 Homo sapiens tumor necrosis factor receptor superfamily, member 25 (TNFF
 NM_148970 Homo sapiens tumor necrosis factor receptor superfamily, member 25 (TNFF
 NM_148971 Homo sapiens tumor necrosis factor receptor superfamily, member 25 (TNFF
 NM_148972 Homo sapiens tumor necrosis factor receptor superfamily, member 25 (TNFF
 NM_148973 Homo sapiens tumor necrosis factor receptor superfamily, member 25 (TNFF
 NM_148974 Homo sapiens tumor necrosis factor receptor superfamily, member 25 (TNFF
 NM_148975 Homo sapiens membrane-spanning 4-domains, subfamily A, member 4 (MS-
 NM_148976 Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 1 (PS)
 NM_148977 Homo sapiens pantothenate kinase 1 (PANK1), transcript variant alpha, mRNA
 NM_148978 Homo sapiens pantothenate kinase 1 (PANK1), transcript variant beta, mRNA
 NM_148979 Homo sapiens cathepsin H (CTSH), transcript variant 2, mRNA
 NM_148980 Homo sapiens Williams Beuren syndrome chromosome region 20C (WBSCF
 NM_149379 Homo sapiens Williams Beuren syndrome chromosome region 20C (WBSCF
 NM_152132 Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 3 (PS)
 NM_152133 Homo sapiens T-cell activation GTPase activating protein (TAGAP), transcrip
 NM_152219 Homo sapiens gap junction protein, chi 1, 31.9kDa (connexin 31.9) (GJC1), r
 NM_152221 Homo sapiens casein kinase 1, epsilon (CSNK1E), transcript variant 1, mRNA
 NM_152222 Homo sapiens tumor necrosis factor receptor superfamily, member 19-like 1 (T
 NM_152223 Homo sapiens protein phosphatase, EF hand calcium-binding domain 1 (PPI
 NM_152224 Homo sapiens protein phosphatase, EF hand calcium-binding domain 1 (PPI
 NM_152225 Homo sapiens protein phosphatase, EF hand calcium-binding domain 1 (PPI
 NM_152226 Homo sapiens protein phosphatase, EF hand calcium-binding domain 1 (PPI
 NM_152227 Homo sapiens sorting nexin 5 (SNX5), transcript variant 1, mRNA
 NM_152230 Homo sapiens inositol polyphosphate multikinase (IPMK), mRNA
 NM_152232 Homo sapiens taste receptor, type 1, member 2 (TAS1R2), mRNA
 NM_152233 Homo sapiens sorting nexin 6 (SNX6), transcript variant 2, mRNA
 NM_152235 Homo sapiens splicing factor, arginine/serine-rich 8 (suppressor-of-white-apr
 NM_152236 Homo sapiens growth arrest-specific 2 like 1 (GAS2L1), transcript variant 2, i
 NM_152237 Homo sapiens growth arrest-specific 2 like 1 (GAS2L1), transcript variant 3, i
 NM_152238 Homo sapiens sorting nexin 7 (SNX7), transcript variant 2, mRNA
 NM_152240 Homo sapiens p53 target zinc finger protein (WIG1), transcript variant 2, mR
 NM_152243 Homo sapiens CDC42 effector protein (Rho GTPase binding) 1 (CDC42EP1)
 NM_152244 Homo sapiens sorting nexin 11 (SNX11), transcript variant 1, mRNA
 NM_152245 Homo sapiens carnitine palmitoyltransferase 1B (muscle) (CPT1B), nuclear c
 NM_152246 Homo sapiens carnitine palmitoyltransferase 1B (muscle) (CPT1B), nuclear c
 NM_152247 Homo sapiens carnitine palmitoyltransferase 1B (muscle) (CPT1B), nuclear c
 NM_152250 Homo sapiens defensin, beta 105 (DEFB105), mRNA
 NM_152251 Homo sapiens defensin, beta 106 (DEFB106), mRNA
 NM_152253 Homo sapiens choline kinase beta (CHKB), transcript variant 2, mRNA
 NM_152255 Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7 (PS)
 NM_152257 Homo sapiens KIAA0889 protein (KIAA0889), mRNA
 NM_152259 Homo sapiens leucine-rich repeat kinase 1 (MGC45866), mRNA
 NM_152260 Homo sapiens chromosome 15 open reading frame 19 (C15orf19), mRNA
 NM_152261 Homo sapiens hypothetical protein MGC17943 (MGC17943), mRNA
 NM_152262 Homo sapiens zinc finger protein 439 (ZNF439), mRNA
 NM_152263 Homo sapiens tropomyosin 3 (TPM3), mRNA
 NM_152264 Homo sapiens solute carrier family 39 (zinc transporter), member 13 (SLC39
 NM_152266 Homo sapiens hypothetical protein MGC32020 (MGC32020), mRNA
 NM_152267 Homo sapiens hypothetical protein FLJ38628 (FLJ38628), mRNA
 NM_152268 Homo sapiens similar to tRNA synthetase class II (DKFZp727A071), mRNA
 NM_152269 Homo sapiens hypothetical protein FLJ38663 (FLJ38663), mRNA
 NM_152270 Homo sapiens hypothetical protein FLJ34922 (FLJ34922), mRNA
 NM_152271 Homo sapiens hypothetical protein FLJ23749 (FLJ23749), mRNA

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NM_152272 Homo sapiens hypothetical protein MGC29816 (MGC29816), mRNA
 NM_152274 Homo sapiens hypothetical protein MGC29729 (MGC29729), mRNA
 NM_152275 Homo sapiens hypothetical protein FLJ13946 (FLJ13946), mRNA
 NM_152277 Homo sapiens dendritic cell-derived ubiquitin-like protein (DC-Ubp), mRNA
 NM_152278 Homo sapiens hypothetical protein MGC23947 (MGC23947), mRNA
 NM_152279 Homo sapiens zinc finger protein 585B (ZNF585B), mRNA
 NM_152280 Homo sapiens synaptotagmin XI (SYT11), mRNA
 NM_152281 Homo sapiens NTKL-binding protein 1 (FLJ11752), mRNA
 NM_152282 Homo sapiens hypothetical protein FLJ23751 (FLJ23751), mRNA
 NM_152283 Homo sapiens zinc finger protein 62 homolog (mouse) (ZFP62), mRNA
 NM_152284 Homo sapiens Snf7 homologue associated with Alix 3 (Shax3), mRNA
 NM_152285 Homo sapiens arrestin domain containing 1 (ARRDC1), mRNA
 NM_152286 Homo sapiens chromosome 9 open reading frame 111 (C9orf111), mRNA
 NM_152287 Homo sapiens zinc finger protein 276 homolog (mouse) (ZFP276), mRNA
 NM_152288 Homo sapiens hypothetical protein MGC13024 (MGC13024), mRNA
 NM_152289 Homo sapiens zinc finger protein 561 (ZNF561), mRNA
 NM_152290 Homo sapiens hypothetical protein MGC35194 (MGC35194), mRNA
 NM_152291 Homo sapiens mucin 7, salivary (MUC7), mRNA
 NM_152292 Homo sapiens RNA (guanine-9-) methyltransferase domain containing 2 (RC
 NM_152295 Homo sapiens threonyl-tRNA synthetase (TARS), mRNA
 NM_152296 Homo sapiens ATPase, Na⁺/K⁺ transporting, alpha 3 polypeptide (ATP1A3),
 NM_152298 Homo sapiens nuclear autoantigenic sperm protein (histone-binding) (NASP)
 NM_152299 Homo sapiens hypothetical protein 384D8_6 (384D8-2), mRNA
 NM_152300 Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 52 (DDX52), transc
 NM_152301 Homo sapiens PP784 protein (PP784), transcript variant 1, mRNA
 NM_152302 Homo sapiens chromosome 20 open reading frame 158 (C20orf158), mRNA
 NM_152303 Homo sapiens zinc finger protein 554 (ZNF554), mRNA
 NM_152304 Homo sapiens hypothetical protein MGC45806 (MGC45806), mRNA
 NM_152305 Homo sapiens x 010 protein (MDS010), mRNA
 NM_152306 Homo sapiens ubiquitin-like, containing PHD and RING finger domains, 2 (U
 NM_152307 Homo sapiens hypothetical protein FLJ40452 (FLJ40452), mRNA
 NM_152308 Homo sapiens hypothetical protein MGC24665 (MGC24665), mRNA
 NM_152309 Homo sapiens phosphoinositide-3-kinase adaptor protein 1 (PIK3AP1), mRN
 NM_152310 Homo sapiens elongation of very long chain fatty acids (FEN1/Elo2, SUR4/E
 NM_152311 Homo sapiens hypothetical protein MGC32871 (MGC32871), mRNA
 NM_152312 Homo sapiens glycosyltransferase-like 1B (GYLTL1B), mRNA
 NM_152313 Homo sapiens solute carrier family 36 (proton/amino acid symporter), membr
 NM_152314 Homo sapiens hypothetical protein MGC34830 (MGC34830), mRNA
 NM_152315 Homo sapiens hypothetical protein MGC34290 (MGC34290), mRNA
 NM_152316 Homo sapiens hypothetical protein FLJ38968 (FLJ38968), mRNA
 NM_152317 Homo sapiens DEP domain containing 4 (DEPDC4), mRNA
 NM_152318 Homo sapiens hypothetical protein MGC40397 (MGC40397), mRNA
 NM_152319 Homo sapiens hypothetical protein MGC35033 (MGC35033), mRNA
 NM_152320 Homo sapiens hypothetical protein FLJ31295 (FLJ31295), mRNA
 NM_152321 Homo sapiens hypothetical protein FLJ32115 (FLJ32115), mRNA
 NM_152322 Homo sapiens BTB (POZ) domain containing 11 (BTBD11), mRNA
 NM_152323 Homo sapiens Spi-C transcription factor (Spi-1/PU.1 related) (SPIC), mRNA
 NM_152324 Homo sapiens hypothetical protein MGC35169 (MGC35169), mRNA
 NM_152325 Homo sapiens hypothetical protein MGC40178 (MGC40178), mRNA
 NM_152326 Homo sapiens ankyrin repeat domain 9 (ANKRD9), mRNA
 NM_152327 Homo sapiens adenylylate kinase 7 (AK7), mRNA
 NM_152328 Homo sapiens adenylosuccinate synthase like 1 (ADSSL1), transcript variant
 NM_152329 Homo sapiens peptidylprolyl isomerase (cyclophilin) like 5 (PPIL5), transcript
 NM_152330 Homo sapiens chromosome 14 open reading frame 31 (C14orf31), mRNA
 NM_152331 Homo sapiens peroxisomal acyl-CoA thioesterase 2B (PTE2B), mRNA
 NM_152332 Homo sapiens membrane targeting (tandem) C2 domain containing 1 (MTAC
 NM_152333 Homo sapiens solute carrier family 25, member 29 (SLC25A29), mRNA

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NM_152334 Homo sapiens FLJ25005 protein (FLJ25005), mRNA
NM_152335 Homo sapiens chromosome 15 open reading frame 27 (C15orf27), mRNA
NM_152336 Homo sapiens hypothetical protein FLJ32310 (FLJ32310), mRNA
NM_152337 Homo sapiens hypothetical protein FLJ32702 (FLJ32702), mRNA
NM_152338 Homo sapiens zymogen granule protein 16 (ZG16), mRNA
NM_152339 Homo sapiens hypothetical protein MGC26885 (MGC26885), mRNA
NM_152340 Homo sapiens hypothetical protein FLJ39075 (FLJ39075), mRNA
NM_152341 Homo sapiens progesterin and adipoQ receptor family member IV (PAQR4), m
NM_152342 Homo sapiens chromodomain protein, Y-like 2 (CDYL2), mRNA
NM_152343 Homo sapiens hypothetical protein FLJ25414 (FLJ25414), mRNA
NM_152344 Homo sapiens hypothetical protein FLJ30656 (FLJ30656), mRNA
NM_152345 Homo sapiens hypothetical protein FLJ25555 (FLJ25555), mRNA
NM_152346 Homo sapiens hypothetical protein MGC34680 (MGC34680), mRNA
NM_152347 Homo sapiens hypothetical protein FLJ40342 (FLJ40342), mRNA
NM_152348 Homo sapiens hypothetical protein FLJ33817 (FLJ33817), mRNA
NM_152349 Homo sapiens hypothetical protein MGC45562 (MGC45562), mRNA
NM_152350 Homo sapiens hypothetical protein MGC40157 (MGC40157), mRNA
NM_152351 Homo sapiens solute carrier family 5 (sodium/glucose cotransporter), memb
NM_152352 Homo sapiens chromosome 18 open reading frame 19 (C18orf19), mRNA
NM_152353 Homo sapiens hypothetical protein MGC33839 (MGC33839), mRNA
NM_152354 Homo sapiens zinc finger protein 285 (ZNF285), mRNA
NM_152355 Homo sapiens zinc finger protein 441 (ZNF441), mRNA
NM_152356 Homo sapiens zinc finger protein 491 (ZNF491), mRNA
NM_152357 Homo sapiens zinc finger protein 440 (ZNF440), mRNA
NM_152358 Homo sapiens hypothetical protein MGC33947 (MGC33947), mRNA
NM_152359 Homo sapiens camitine palmitoyltransferase 1C (CPT1C), mRNA
NM_152360 Homo sapiens zinc finger protein 573 (ZNF573), mRNA
NM_152361 Homo sapiens hypothetical protein FLJ38944 (FLJ38944), mRNA
NM_152362 Homo sapiens hypothetical protein MGC17791 (MGC17791), mRNA
NM_152363 Homo sapiens hypothetical protein FLJ39369 (FLJ39369), mRNA
NM_152365 Homo sapiens hypothetical protein FLJ34633 (FLJ34633), mRNA
NM_152366 Homo sapiens hypothetical protein MGC33338 (MGC33338), mRNA
NM_152367 Homo sapiens hypothetical protein FLJ38716 (FLJ38716), mRNA
NM_152369 Homo sapiens hypothetical protein MGC45474 (MGC45474), mRNA
NM_152371 Homo sapiens hypothetical protein MGC26818 (MGC26818), mRNA
NM_152372 Homo sapiens myomesin family, member 3 (MYOM3), mRNA
NM_152373 Homo sapiens hypothetical protein MGC27466 (MGC27466), mRNA
NM_152374 Homo sapiens hypothetical protein FLJ38984 (FLJ38984), mRNA
NM_152375 Homo sapiens hypothetical protein FLJ38753 (FLJ38753), mRNA
NM_152376 Homo sapiens UBX domain containing 3 (UBXD3), mRNA
NM_152377 Homo sapiens hypothetical protein MGC34837 (MGC34837), mRNA
NM_152378 Homo sapiens hypothetical protein FLJ31052 (FLJ31052), mRNA
NM_152379 Homo sapiens hypothetical protein DKFZp547B1713 (DKFZp547B1713), mF
NM_152382 Homo sapiens hypothetical protein FLJ37953 (FLJ37953), mRNA
NM_152383 Homo sapiens hypothetical protein MGC42174 (MGC42174), mRNA
NM_152384 Homo sapiens Bardet-Biedl syndrome 5 (BBS5), mRNA
NM_152385 Homo sapiens hypothetical protein FLJ31438 (FLJ31438), mRNA
NM_152386 Homo sapiens sphingosine-1-phosphate phosphatase 2 (SGPP2), mRNA
NM_152387 Homo sapiens hypothetical protein FLJ31322 (FLJ31322), mRNA
NM_152388 Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region,
NM_152389 Homo sapiens hypothetical protein MGC35338 (MGC35338), mRNA
NM_152390 Homo sapiens hypothetical protein MGC33926 (MGC33926), mRNA
NM_152391 Homo sapiens chromosome 2 open reading frame 22 (C2orf22), mRNA
NM_152392 Homo sapiens AHA1, activator of heat shock 90kDa protein ATPase homolo
NM_152393 Homo sapiens kelch repeat and BTB (POZ) domain containing 5 (KBTBD5),
NM_152394 Homo sapiens hypothetical protein MGC39662 (MGC39662), mRNA
NM_152395 Homo sapiens hypothetical protein FLJ31265 (FLJ31265), mRNA

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NM_152396 Homo sapiens hypothetical protein MGC24132 (MGC24132), mRNA
 NM_152397 Homo sapiens hypothetical protein MGC39725 (MGC39725), mRNA
 NM_152398 Homo sapiens hypothetical protein MGC45416 (MGC45416), mRNA
 NM_152399 Homo sapiens hypothetical protein FLJ30834 (FLJ30834), mRNA
 NM_152400 Homo sapiens hypothetical protein FLJ39370 (FLJ39370), mRNA
 NM_152401 Homo sapiens phosphatase-like 2 (PDCD2), mRNA
 NM_152402 Homo sapiens translocation associated membrane protein 1-like 1 (TRAM1L)
 NM_152403 Homo sapiens hypothetical protein FLJ39155 (FLJ39155), transcript variant
 NM_152404 Homo sapiens hypothetical protein FLJ34658 (FLJ34658), mRNA
 NM_152405 Homo sapiens junction-mediating and regulatory protein (JMY), mRNA
 NM_152407 Homo sapiens GrpE-like 2, mitochondrial (E. coli) (GRPEL2), mRNA
 NM_152408 Homo sapiens hypothetical protein FLJ35779 (FLJ35779), mRNA
 NM_152409 Homo sapiens hypothetical protein FLJ37562 (FLJ37562), mRNA
 NM_152410 Homo sapiens PARK2 co-regulated (PACRG), mRNA
 NM_152411 Homo sapiens hypothetical protein DKFZp762I137 (DKFZp762I137), mRNA
 NM_152412 Homo sapiens zinc finger protein 572 (ZNF572), mRNA
 NM_152413 Homo sapiens hypothetical protein MGC33309 (MGC33309), mRNA
 NM_152414 Homo sapiens basic helix-loop-helix domain containing, class B, 5 (BHLHB5)
 NM_152415 Homo sapiens hepatocellular carcinoma related protein 1 (FLJ32642), mRNA
 NM_152416 Homo sapiens hypothetical protein MGC40214 (MGC40214), mRNA
 NM_152417 Homo sapiens hypothetical protein FLJ32370 (FLJ32370), mRNA
 NM_152418 Homo sapiens hypothetical protein FLJ35775 (FLJ35775), mRNA
 NM_152420 Homo sapiens chromosome 9 open reading frame 41 (C9orf41), mRNA
 NM_152421 Homo sapiens hypothetical protein MGC20262 (MGC20262), mRNA
 NM_152422 Homo sapiens protein tyrosine phosphatase domain containing 1 (PTPDC1),
 NM_152423 Homo sapiens hypothetical protein FLJ33516 (FLJ33516), mRNA
 NM_152424 Homo sapiens hypothetical protein FLJ39827 (FLJ39827), mRNA
 NM_152425 Homo sapiens hypothetical protein FLJ40249 (FLJ40249), mRNA
 NM_152427 Homo sapiens cofilin pseudogene 1 (CFLP1), mRNA
 NM_152428 Homo sapiens FERM and PDZ domain containing 2 (FRMPD2), mRNA
 NM_152429 Homo sapiens chromosome 10 open reading frame 13 (C10orf13), mRNA
 NM_152430 Homo sapiens hypothetical protein MGC24137 (MGC24137), mRNA
 NM_152431 Homo sapiens pili-like 4 (Drosophila) (PIWIL4), mRNA
 NM_152433 Homo sapiens kelch repeat and BTB (POZ) domain containing 3 (KBTBD3),
 NM_152434 Homo sapiens CWF19-like 2, cell cycle c-control (S. pombe) (CWF19L2), mRNA
 NM_152435 Homo sapiens hypothetical protein MGC35366 (MGC35366), mRNA
 NM_152436 Homo sapiens hypothetical protein MGC39497 (MGC39497), mRNA
 NM_152437 Homo sapiens hypothetical protein DKFZp761B128 (DKFZp761B128), mRNA
 NM_152439 Homo sapiens vitelliform macular dystrophy 2-like 3 (VMD2L3), mRNA
 NM_152440 Homo sapiens hypothetical protein FLJ32549 (FLJ32549), mRNA
 NM_152441 Homo sapiens F-box and leucine-rich repeat protein 14 (FBXL14), mRNA
 NM_152442 Homo sapiens RAD9 homolog B (S. cerevisiae) (RAD9B), mRNA
 NM_152443 Homo sapiens retinol dehydrogenase 12 (all-trans and 9-cis) (RDH12), mRNA
 NM_152444 Homo sapiens zinc binding alcohol dehydrogenase, domain containing 1 (ZA)
 NM_152445 Homo sapiens chromosome 14 open reading frame 44 (C14orf44), mRNA
 NM_152447 Homo sapiens leucine rich repeat and fibronectin type III domain containing 1
 NM_152448 Homo sapiens hypothetical protein MGC33951 (MGC33951), mRNA
 NM_152449 Homo sapiens hypothetical protein FLJ33008 (FLJ33008), mRNA
 NM_152450 Homo sapiens hypothetical protein MGC26690 (MGC26690), mRNA
 NM_152451 Homo sapiens GRINL1A complex upstream protein (Gup1), mRNA
 NM_152453 Homo sapiens hypothetical protein MGC35118 (MGC35118), mRNA
 NM_152454 Homo sapiens hypothetical protein FLJ31461 (FLJ31461), mRNA
 NM_152455 Homo sapiens hypothetical protein FLJ35867 (FLJ35867), mRNA
 NM_152456 Homo sapiens hypothetical protein MGC34647 (MGC34647), mRNA
 NM_152457 Homo sapiens zinc finger protein 597 (ZNF597), mRNA
 NM_152458 Homo sapiens hypothetical protein FLJ32130 (FLJ32130), mRNA
 NM_152459 Homo sapiens hypothetical protein MGC45438 (MGC45438), mRNA

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NM_152460 Homo sapiens hypothetical protein FLJ31882 (FLJ31882), mRNA
 NM_152461 Homo sapiens ER to nucleus signalling 1 (ERN1), transcript variant 2, mRNA
 NM_152462 Homo sapiens transmembrane protein 21A (TMEM21A), mRNA
 NM_152463 Homo sapiens essential meiotic endonuclease 1 homolog 1 (S. pombe) (EMI
 NM_152464 Homo sapiens chromosome 17 open reading frame 32 (C17orf32), mRNA
 NM_152465 Homo sapiens hypothetical protein MGC39650 (MGC39650), mRNA
 NM_152466 Homo sapiens hypothetical protein FLJ25168 (FLJ25168), mRNA
 NM_152467 Homo sapiens kelch-like 10 (Drosophila) (KLHL10), mRNA
 NM_152468 Homo sapiens epidermodysplasia verruciformis 2 (EVER2), mRNA
 NM_152470 Homo sapiens chromosome 18 open reading frame 23 (C18orf23), mRNA
 NM_152472 Homo sapiens zinc finger protein 578 (ZNF578), mRNA
 NM_152473 Homo sapiens hypothetical protein FLJ32214 (FLJ32214), mRNA
 NM_152474 Homo sapiens chromosome 19 open reading frame 18 (C19orf18), mRNA
 NM_152475 Homo sapiens hypothetical protein MGC34079 (MGC34079), mRNA
 NM_152476 Homo sapiens zinc finger protein 560 (ZNF560), mRNA
 NM_152477 Homo sapiens zinc finger protein 565 (ZNF565), mRNA
 NM_152478 Homo sapiens zinc finger protein 583 (ZNF583), mRNA
 NM_152479 Homo sapiens hypothetical protein MGC33962 (MGC33962), mRNA
 NM_152480 Homo sapiens chromosome 19 open reading frame 23 (C19orf23), mRNA
 NM_152481 Homo sapiens hypothetical protein FLJ25660 (FLJ25660), mRNA
 NM_152482 Homo sapiens chromosome 19 open reading frame 25 (C19orf25), mRNA
 NM_152483 Homo sapiens hypothetical protein FLJ25328 (FLJ25328), mRNA
 NM_152484 Homo sapiens zinc finger protein 569 (ZNF569), mRNA
 NM_152485 Homo sapiens hypothetical protein FLJ25078 (FLJ25078), mRNA
 NM_152486 Homo sapiens sterile alpha motif domain containing 11 (SAMD11), mRNA
 NM_152487 Homo sapiens hypothetical protein FLJ31842 (FLJ31842), mRNA
 NM_152488 Homo sapiens hypothetical protein FLJ32833 (FLJ32833), mRNA
 NM_152489 Homo sapiens hypothetical protein MGC35130 (MGC35130), mRNA
 NM_152490 Homo sapiens beta 1,3-N-acetylgalactosaminyltransferase-II (MGC39558), n
 NM_152491 Homo sapiens hypothetical protein FLJ32569 (FLJ32569), mRNA
 NM_152492 Homo sapiens hypothetical protein FLJ32825 (FLJ32825), mRNA
 NM_152493 Homo sapiens FLJ25476 protein (FLJ25476), mRNA
 NM_152494 Homo sapiens hypothetical protein FLJ32785 (FLJ32785), mRNA
 NM_152495 Homo sapiens hypothetical protein FLJ38993 (FLJ38993), mRNA
 NM_152496 Homo sapiens hypothetical protein FLJ31434 (FLJ31434), mRNA
 NM_152497 Homo sapiens hypothetical protein FLJ32208 (FLJ32208), mRNA
 NM_152498 Homo sapiens hypothetical protein FLJ32000 (FLJ32000), mRNA
 NM_152499 Homo sapiens hypothetical protein MGC45441 (MGC45441), mRNA
 NM_152500 Homo sapiens hypothetical protein FLJ33084 (FLJ33084), mRNA
 NM_152501 Homo sapiens interferon-inducible protein X (IFI), transcript variant a1, mR
 NM_152503 Homo sapiens chromosome 20 open reading frame 132 (C20orf132), transcr
 NM_152504 Homo sapiens hypothetical protein FLJ25067 (FLJ25067), mRNA
 NM_152505 Homo sapiens chromosome 21 open reading frame 13 (C21orf13), mRNA
 NM_152506 Homo sapiens chromosome 21 open reading frame 129 (C21orf129), mRNA
 NM_152507 Homo sapiens chromosome 21 open reading frame 128 (C21orf128), mRNA
 NM_152509 Homo sapiens hypothetical protein FLJ31568 (FLJ31568), mRNA
 NM_152510 Homo sapiens hypothetical protein MGC26710 (MGC26710), mRNA
 NM_152511 Homo sapiens dual specificity phosphatase 18 (DUSP18), mRNA
 NM_152512 Homo sapiens hypothetical protein FLJ25421 (FLJ25421), mRNA
 NM_152515 Homo sapiens hypothetical protein FLJ40629 (FLJ40629), mRNA
 NM_152516 Homo sapiens copper metabolism (Murr1) domain containing 1 (COMMD1),
 NM_152517 Homo sapiens hypothetical protein FLJ30990 (FLJ30990), mRNA
 NM_152519 Homo sapiens hypothetical protein FLJ23861 (FLJ23861), mRNA
 NM_152520 Homo sapiens zinc finger protein 533 (ZNF533), mRNA
 NM_152522 Homo sapiens ADP-ribosylation-like factor 6-interacting protein 6 (MGC3386
 NM_152523 Homo sapiens hypothetical protein FLJ40432 (FLJ40432), mRNA
 NM_152524 Homo sapiens shugoshin-like 2 (S. pombe) (SGOL2), mRNA

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NM_152525 Homo sapiens hypothetical protein FLJ25351 (FLJ25351), mRNA
 NM_152526 Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region,
 NM_152527 Homo sapiens solute carrier family 16 (monocarboxylic acid transporters), m
 NM_152528 Homo sapiens WD repeat and SAM domain containing 1 (WDSAM1), mRNA
 NM_152529 Homo sapiens G protein-coupled receptor 155 (GPR155), mRNA
 NM_152531 Homo sapiens hypothetical protein FLJ35155 (FLJ35155), mRNA
 NM_152533 Homo sapiens hypothetical protein MGC34728 (MGC34728), mRNA
 NM_152534 Homo sapiens hypothetical protein FLJ32685 (FLJ32685), mRNA
 NM_152536 Homo sapiens FYVE, RhoGEF and PH domain containing 5 (FGD5), mRNA
 NM_152538 Homo sapiens immunoglobulin superfamily, member 11 (IGSF11), mRNA
 NM_152539 Homo sapiens hypothetical protein FLJ32859 (FLJ32859), mRNA
 NM_152540 Homo sapiens sec1 family domain containing 2 (SCFD2), mRNA
 NM_152542 Homo sapiens hypothetical protein DKFZp761G058 (DKFZp761G058), mRN
 NM_152543 Homo sapiens hypothetical protein FLJ25371 (FLJ25371), mRNA
 NM_152544 Homo sapiens hypothetical protein FLJ35725 (FLJ35725), mRNA
 NM_152545 Homo sapiens RasGEF domain family, member 1B (RASGEF1B), mRNA
 NM_152546 Homo sapiens hypothetical protein FLJ25286 (FLJ25286), mRNA
 NM_152547 Homo sapiens butyrophilin-like 9 (BTNL9), mRNA
 NM_152548 Homo sapiens hypothetical protein FLJ25333 (FLJ25333), mRNA
 NM_152549 Homo sapiens hypothetical protein MGC39633 (MGC39633), mRNA
 NM_152550 Homo sapiens SH3 domain containing ring finger 2 (SH3RF2), mRNA
 NM_152551 Homo sapiens chromosome 6 open reading frame 151 (C6orf151), mRNA
 NM_152552 Homo sapiens sterile alpha motif domain containing 3 (SAMD3), mRNA
 NM_152553 Homo sapiens IBR domain containing 1 (IBRDC1), mRNA
 NM_152554 Homo sapiens chromosome 6 open reading frame 195 (C6orf195), mRNA
 NM_152556 Homo sapiens hypothetical protein FLJ31818 (FLJ31818), mRNA
 NM_152557 Homo sapiens hypothetical protein FLJ31413 (FLJ31413), mRNA
 NM_152558 Homo sapiens KIAA1023 protein (KIAA1023), mRNA
 NM_152559 Homo sapiens Williams Beuren syndrome chromosome region 27 (WBSR2
 NM_152562 Homo sapiens cell division cycle associated 2 (CDCA2), mRNA
 NM_152563 Homo sapiens hypothetical protein FLJ10661 (FLJ10661), mRNA
 NM_152564 Homo sapiens Cohen syndrome 1 (COH1), transcript variant 1, mRNA
 NM_152565 Homo sapiens ATPase, H⁺ transporting, lysosomal 38kDa, V0 subunit d iso
 NM_152568 Homo sapiens hypothetical protein FLJ25169 (FLJ25169), mRNA
 NM_152569 Homo sapiens chromosome 9 open reading frame 66 (C9orf66), mRNA
 NM_152570 Homo sapiens hypothetical protein FLJ31810 (FLJ31810), mRNA
 NM_152571 Homo sapiens hypothetical protein FLJ36779 (FLJ36779), mRNA
 NM_152572 Homo sapiens chromosome 9 open reading frame 98 (C9orf98), mRNA
 NM_152573 Homo sapiens RAS and EF hand domain containing (RASEF), mRNA
 NM_152574 Homo sapiens chromosome 9 open reading frame 52 (C9orf52), mRNA
 NM_152577 Homo sapiens hypothetical protein FLJ25735 (FLJ25735), mRNA
 NM_152578 Homo sapiens fragile X mental retardation 1 neighbor (FMR1NB), mRNA
 NM_152579 Homo sapiens hypothetical protein FLJ38564 (FLJ38564), mRNA
 NM_152581 Homo sapiens motile sperm domain containing 2 (MOSPD2), mRNA
 NM_152582 Homo sapiens hypothetical protein MGC27005 (MGC27005), mRNA
 NM_152583 Homo sapiens hypothetical protein MGC40053 (MGC40053), mRNA
 NM_152584 Homo sapiens heat shock transcription factor, Y-linked 1 (HSFY1), transcript
 NM_152585 Homo sapiens RNA binding motif protein, Y-linked, family 1 (MGC33094), m
 NM_152586 Homo sapiens ubiquitin specific protease 54 (USP54), mRNA
 NM_152587 Homo sapiens hypothetical protein MGC33948 (MGC33948), mRNA
 NM_152588 Homo sapiens hypothetical protein DKFZp762A217 (DKFZp762A217), mRN
 NM_152589 Homo sapiens hypothetical protein FLJ35821 (FLJ35821), mRNA
 NM_152590 Homo sapiens hypothetical protein FLJ30004 (FLJ30004), mRNA
 NM_152591 Homo sapiens hypothetical protein FLJ35843 (FLJ35843), mRNA
 NM_152592 Homo sapiens chromosome 14 open reading frame 49 (C14orf49), mRNA
 NM_152594 Homo sapiens sprouty-related, EVH1 domain containing 1 (SPRED1), mRN
 NM_152595 Homo sapiens piggyBac transposable element derived 4 (PGBD4), mRNA

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NM_152586 Homo sapiens hypothetical protein MGC33637 (MGC33637), mRNA
 NM_152587 Homo sapiens fibrous sheath interacting protein 1 (FSIP1), mRNA
 NM_152588 Homo sapiens hypothetical protein FLJ35757 (FLJ35757), mRNA
 NM_152589 Homo sapiens hypothetical protein FLJ35773 (FLJ35773), mRNA
 NM_152600 Homo sapiens zinc finger protein 579 (ZNF579), mRNA
 NM_152601 Homo sapiens hypothetical protein FLJ38281 (FLJ38281), mRNA
 NM_152602 Homo sapiens zinc finger protein 433 (ZNF433), mRNA
 NM_152603 Homo sapiens zinc finger protein 567 (ZNF567), mRNA
 NM_152604 Homo sapiens zinc finger protein 383 (ZNF383), mRNA
 NM_152605 Homo sapiens hypothetical protein FLJ37549 (FLJ37549), mRNA
 NM_152606 Homo sapiens zinc finger protein 540 (ZNF540), mRNA
 NM_152607 Homo sapiens hypothetical protein FLJ40201 (FLJ40201), mRNA
 NM_152608 Homo sapiens hypothetical protein FLJ35382 (FLJ35382), mRNA
 NM_152609 Homo sapiens hypothetical protein FLJ32001 (FLJ32001), mRNA
 NM_152610 Homo sapiens hypothetical protein FLJ35728 (FLJ35728), mRNA
 NM_152611 Homo sapiens chromosome 20 open reading frame 75 (C20orf75), mRNA
 NM_152612 Homo sapiens hypothetical protein FLJ38046 (FLJ38046), mRNA
 NM_152613 Homo sapiens hypothetical protein MGC26816 (MGC26816), mRNA
 NM_152614 Homo sapiens hypothetical protein MGC35154 (MGC35154), mRNA
 NM_152615 Homo sapiens hypothetical protein FLJ40597 (FLJ40597), mRNA
 NM_152616 Homo sapiens tripartite motif-containing 42 (TRIM42), mRNA
 NM_152617 Homo sapiens hypothetical protein FLJ35794 (FLJ35794), mRNA
 NM_152618 Homo sapiens hypothetical protein FLJ35630 (FLJ35630), mRNA
 NM_152619 Homo sapiens hypothetical protein MGC45428 (MGC45428), mRNA
 NM_152620 Homo sapiens ring finger protein 129 (RNF129), mRNA
 NM_152621 Homo sapiens hypothetical protein MGC26963 (MGC26963), mRNA
 NM_152622 Homo sapiens hypothetical protein FLJ35954 (FLJ35954), mRNA
 NM_152623 Homo sapiens CDC20-like protein (FLJ37927), mRNA
 NM_152624 Homo sapiens decapping enzyme hDcp2 (DCP2), mRNA
 NM_152625 Homo sapiens zinc finger protein 366 (ZNF366), mRNA
 NM_152626 Homo sapiens zinc finger protein 92 (HTF12) (ZNF92), mRNA
 NM_152628 Homo sapiens hypothetical protein MGC39715 (MGC39715), mRNA
 NM_152629 Homo sapiens GLIS family zinc finger 3 (GLIS3), mRNA
 NM_152630 Homo sapiens hypothetical protein MGC26999 (MGC26999), mRNA
 NM_152631 Homo sapiens hypothetical protein FLJ35782 (FLJ35782), mRNA
 NM_152632 Homo sapiens chromosome X open reading frame 22 (CXorf22), mRNA
 NM_152633 Homo sapiens hypothetical protein FLJ34064 (FLJ34064), mRNA
 NM_152635 Homo sapiens oncoprotein induced transcript 3 (OIT3), mRNA
 NM_152636 Homo sapiens hypothetical protein FLJ33979 (FLJ33979), mRNA
 NM_152637 Homo sapiens hypothetical protein MGC17301 (MGC17301), mRNA
 NM_152638 Homo sapiens hypothetical protein MGC26598 (MGC26598), mRNA
 NM_152640 Homo sapiens decapping enzyme hDcp1b (DCP1B), mRNA
 NM_152643 Homo sapiens kinase non-catalytic C-lobe domain (KIND) containing 1 (KND
 NM_152644 Homo sapiens family with sequence similarity 24, member B (FAM24B), mR
 NM_152647 Homo sapiens hypothetical protein FLJ32800 (FLJ32800), mRNA
 NM_152649 Homo sapiens hypothetical protein FLJ34389 (FLJ34389), mRNA
 NM_152652 Homo sapiens zinc finger protein 553 (ZNF553), mRNA
 NM_152653 Homo sapiens ubiquitin-conjugating enzyme E2E 2 (UBC4/5 homolog, yeast
 NM_152654 Homo sapiens hypothetical protein FLJ38607 (FLJ38607), mRNA
 NM_152655 Homo sapiens zinc finger protein 585A (ZNF585A), transcript variant 1, mRN
 NM_152657 Homo sapiens gametogenetin (GON), transcript variant 1, mRNA
 NM_152658 Homo sapiens THAP domain containing 8 (THAP8), mRNA
 NM_152660 Homo sapiens hypothetical protein MGC34648 (MGC34648), mRNA
 NM_152662 Homo sapiens hypothetical protein FLJ23867 (FLJ23867), mRNA
 NM_152663 Homo sapiens Ral GEF with PH domain and SH3 binding motif 2 (RALGPS2
 NM_152665 Homo sapiens hypothetical protein FLJ40873 (FLJ40873), mRNA
 NM_152666 Homo sapiens hypothetical protein FLJ40773 (FLJ40773), mRNA

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NM_152667 Homo sapiens chromosome 20 open reading frame 147 (C20orf147), mRNA
 NM_152670 Homo sapiens hypothetical protein FLJ25369 (FLJ25369), mRNA
 NM_152671 Homo sapiens phosphatidylinositol-3-phosphate/phosphatidylinositol 5-kinas
 NM_152672 Homo sapiens organic solute transporter alpha (OSTalpha), mRNA
 NM_152673 Homo sapiens mucin 20 (MUC20), mRNA
 NM_152675 Homo sapiens hypothetical protein FLJ23754 (FLJ23754), mRNA
 NM_152676 Homo sapiens F-box protein 15 (FBXO15), mRNA
 NM_152677 Homo sapiens zinc finger protein 494 (ZNF494), mRNA
 NM_152678 Homo sapiens hypothetical protein FLJ34969 (FLJ34969), mRNA
 NM_152679 Homo sapiens solute carrier family 10 (sodium/bile acid cotransporter family)
 NM_152680 Homo sapiens hypothetical protein FLJ32028 (FLJ32028), mRNA
 NM_152681 Homo sapiens hypothetical protein FLJ38482 (FLJ38482), mRNA
 NM_152682 Homo sapiens hypothetical protein MGC10198 (MGC10198), mRNA
 NM_152683 Homo sapiens hypothetical protein FLJ33167 (FLJ33167), mRNA
 NM_152684 Homo sapiens hypothetical protein FLJ39653 (FLJ39653), mRNA
 NM_152685 Homo sapiens solute carrier family 23 (nucleobase transporters), member 1
 NM_152686 Homo sapiens hypothetical protein MGC29463 (MGC29463), mRNA
 NM_152687 Homo sapiens hypothetical protein FLJ33641 (FLJ33641), mRNA
 NM_152688 Homo sapiens KH domain containing, RNA binding, signal transduction asso
 NM_152689 Homo sapiens hypothetical protein MGC9712 (MGC9712), mRNA
 NM_152690 Homo sapiens dolichyl-phosphate mannosyltransferase polypeptide 2, regul
 NM_152692 Homo sapiens core 1 UDP-galactose:4-acetylgalactosamine-alpha-R beta 1
 NM_152693 Homo sapiens hypothetical protein MGC34827 (MGC34827), mRNA
 NM_152694 Homo sapiens zinc finger, CCHC domain containing 5 (ZCCH5), mRNA
 NM_152695 Homo sapiens hypothetical protein FLJ23614 (FLJ23614), mRNA
 NM_152696 Homo sapiens homeodomain interacting protein kinase 1 (HIPK1), transcript
 NM_152697 Homo sapiens hypothetical protein MGC34032 (MGC34032), mRNA
 NM_152698 Homo sapiens hypothetical protein FLJ38377 (FLJ38377), mRNA
 NM_152699 Homo sapiens SUMO1/sentrin specific protease 5 (SENP5), mRNA
 NM_152700 Homo sapiens hypothetical protein MGC26597 (MGC26597), mRNA
 NM_152701 Homo sapiens ATP binding cassette gene, sub-family A (ABC1), member 13
 NM_152702 Homo sapiens chromosome 9 open reading frame 94 (C9orf94), mRNA
 NM_152704 Homo sapiens hypothetical protein FLJ25477 (FLJ25477), transcript variant
 NM_152705 Homo sapiens hypothetical protein MGC9850 (MGC9850), mRNA
 NM_152706 Homo sapiens hypothetical protein MGC28647 (MGC28647), mRNA
 NM_152707 Homo sapiens solute carrier family 25 (mitochondrial carrier; Graves disease
 NM_152710 Homo sapiens chromosome 10 open reading frame 27 (C10orf27), mRNA
 NM_152713 Homo sapiens integral membrane protein 1 (ITM1), mRNA
 NM_152715 Homo sapiens hypothetical protein MGC10233 (MGC10233), mRNA
 NM_152716 Homo sapiens hypothetical protein FLJ36874 (FLJ36874), mRNA
 NM_152717 Homo sapiens hypothetical protein MGC35295 (MGC35295), mRNA
 NM_152718 Homo sapiens hypothetical protein FLJ32009 (FLJ32009), mRNA
 NM_152719 Homo sapiens testis-specific leucine zipper protein nurlt (NURIT), mRNA
 NM_152720 Homo sapiens NIMA (never in mitosis gene a)-related kinase 3 (NEK3), trans
 NM_152721 Homo sapiens docking protein 5-like (DOK5L), mRNA
 NM_152722 Homo sapiens hypothetical protein FLJ25530 (FLJ25530), mRNA
 NM_152723 Homo sapiens hypothetical protein FLJ38159 (FLJ38159), mRNA
 NM_152724 Homo sapiens Ras suppressor protein 1 (RSU1), transcript variant 2, mRNA
 NM_152725 Homo sapiens solute carrier family 39 (zinc transporter), member 12 (SLC39
 NM_152726 Homo sapiens Smhs2 homolog (rat) (FLJ34588), mRNA
 NM_152727 Homo sapiens copine II (CPNE2), mRNA
 NM_152728 Homo sapiens chromosome 18 open reading frame 20 (C18orf20), mRNA
 NM_152729 Homo sapiens 5'-nucleotidase, cytosolic II-like 1 (NT5C2L1), mRNA
 NM_152730 Homo sapiens chromosome 6 open reading frame 170 (C6orf170), mRNA
 NM_152731 Homo sapiens chromosome 6 open reading frame 65 (C6orf65), mRNA
 NM_152732 Homo sapiens chromosome 6 open reading frame 206 (C6orf206), mRNA
 NM_152733 Homo sapiens BTB (POZ) domain containing 9 (BTBD9), mRNA

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NM_152734 Homo sapiens chromosome 6 open reading frame 89 (C6orf89), mRNA
 NM_152735 Homo sapiens zinc finger and BTB domain containing 9 (ZBTB9), mRNA
 NM_152736 Homo sapiens zinc finger protein 187 (ZNF187), mRNA
 NM_152737 Homo sapiens hypothetical protein MGC33993 (MGC33993), mRNA
 NM_152738 Homo sapiens hypothetical protein MGC40222 (MGC40222), mRNA
 NM_152739 Homo sapiens homeo box A9 (HOXA9), transcript variant 1, mRNA
 NM_152740 Homo sapiens 3-hydroxyisobutyrate dehydrogenase (HIBADH), mRNA
 NM_152742 Homo sapiens glypican 2 (cerebroglycan) (GPC2), mRNA
 NM_152743 Homo sapiens chromosome 7 open reading frame 27 (C7orf27), mRNA
 NM_152744 Homo sapiens sidekick homolog 1 (chicken) (SDK1), mRNA
 NM_152745 Homo sapiens neurexophilin 1 (NXPH1), mRNA
 NM_152747 Homo sapiens hypothetical protein DKFpZp586l1420 (DKFpZp586l1420), mRNA
 NM_152748 Homo sapiens hypothetical protein FLJ31340 (FLJ31340), mRNA
 NM_152749 Homo sapiens hypothetical protein MGC33190 (MGC33190), mRNA
 NM_152750 Homo sapiens hypothetical protein FLJ23834 (FLJ23834), mRNA
 NM_152751 Homo sapiens chromosome 10 open reading frame 30 (C10orf30), mRNA
 NM_152753 Homo sapiens signal peptide, CUB domain, EGF-like 3 (SCUBE3), mRNA
 NM_152754 Homo sapiens sema domain, immunoglobulin domain (Ig), short basic domain
 NM_152755 Homo sapiens hypothetical protein MGC40499 (MGC40499), mRNA
 NM_152757 Homo sapiens hypothetical protein FLJ30313 (FLJ30313), mRNA
 NM_152758 Homo sapiens YTH domain family 3 (YTHDF3), mRNA
 NM_152759 Homo sapiens hypothetical protein MGC35140 (MGC35140), mRNA
 NM_152760 Homo sapiens hypothetical protein FLJ30934 (FLJ30934), mRNA
 NM_152761 Homo sapiens hypothetical protein FLJ25444 (FLJ25444), mRNA
 NM_152762 Homo sapiens testis specific, 10 interacting protein (TSGA10IP), mRNA
 NM_152763 Homo sapiens hypothetical protein MGC26989 (MGC26989), mRNA
 NM_152764 Homo sapiens hypothetical protein MGC35212 (MGC35212), mRNA
 NM_152765 Homo sapiens hypothetical protein MGC33510 (MGC33510), mRNA
 NM_152766 Homo sapiens hypothetical protein MGC40107 (MGC40107), mRNA
 NM_152769 Homo sapiens chromosome 19 open reading frame 26 (C19orf26), mRNA
 NM_152770 Homo sapiens hypothetical protein MGC35043 (MGC35043), mRNA
 NM_152771 Homo sapiens chromosome 19 open reading frame 34 (C19orf34), mRNA
 NM_152772 Homo sapiens hypothetical protein MGC40368 (MGC40368), mRNA
 NM_152773 Homo sapiens hypothetical protein MGC33212 (MGC33212), mRNA
 NM_152774 Homo sapiens hypothetical protein MGC42090 (MGC42090), mRNA
 NM_152775 Homo sapiens hypothetical protein MGC33607 (MGC33607), mRNA
 NM_152776 Homo sapiens hypothetical protein MGC40579 (MGC40579), mRNA
 NM_152777 Homo sapiens chromosome 14 open reading frame 48 (C14orf48), mRNA
 NM_152778 Homo sapiens hypothetical protein MGC33302 (MGC33302), mRNA
 NM_152779 Homo sapiens hypothetical protein MGC26856 (MGC26856), mRNA
 NM_152780 Homo sapiens hypothetical protein FLJ14503 (FLJ14503), mRNA
 NM_152781 Homo sapiens hypothetical protein FLJ32830 (FLJ32830), mRNA
 NM_152782 Homo sapiens hypothetical protein MGC33329 (MGC33329), mRNA
 NM_152783 Homo sapiens hypothetical protein MGC25181 (MGC25181), mRNA
 NM_152784 Homo sapiens hypothetical protein MGC39581 (MGC39581), mRNA
 NM_152785 Homo sapiens germinal center expressed transcript 2 (GCET2), mRNA
 NM_152786 Homo sapiens chromosome 9 open reading frame 43 (C9orf43), mRNA
 NM_152787 Homo sapiens TAK1-binding protein 3 (TAB3), transcript variant 1, mRNA
 NM_152788 Homo sapiens E2a-Pbx1-associated protein (EB-1), transcript variant 1, mRNA
 NM_152789 Homo sapiens hypothetical protein MGC40405 (MGC40405), mRNA
 NM_152791 Homo sapiens zinc finger protein 555 (ZNF555), mRNA
 NM_152792 Homo sapiens hypothetical protein FLJ25084 (FLJ25084), mRNA
 NM_152793 Homo sapiens hypothetical protein Ells1 (Ells1), mRNA
 NM_152794 Homo sapiens hypoxia inducible factor 3, alpha subunit (HIF3A), transcript v1
 NM_152795 Homo sapiens hypoxia inducible factor 3, alpha subunit (HIF3A), transcript v1
 NM_152796 Homo sapiens hypoxia inducible factor 3, alpha subunit (HIF3A), transcript v1
 NM_152826 Homo sapiens sorting nexin 1 (SNX1), transcript variant t 3, mRNA

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NM_152827 Homo sapiens sorting nexin 3 (SNX3), transcript variant 2, mRNA
 NM_152828 Homo sapiens sorting nexin 3 (SNX3), transcript variant 3, mRNA
 NM_152829 Homo sapiens testis derived transcript (3 LIM domains) (TES), transcript vari
 NM_152830 Homo sapiens angiotensin I converting enzyme (peptidyl-dipeptidase A) 1 (A
 NM_152831 Homo sapiens angiotensin I converting enzyme (peptidyl-dipeptidase A) 1 (A
 NM_152832 Homo sapiens Mouse Mammary Tumor Virus Receptor homolog 1 (MTVR1)
 NM_152834 Homo sapiens transmembrane protein 18 (TMEM18), mRNA
 NM_152835 Homo sapiens casein kinase (LOC149420), mRNA
 NM_152836 Homo sapiens sorting nexin 16 (SNX16), transcript variant 2, mRNA
 NM_152837 Homo sapiens sorting nexin 16 (SNX16), transcript variant 3, mRNA
 NM_152838 Homo sapiens RNA binding motif protein 12 (RBM12), transcript variant 2, m
 NM_152840 Homo sapiens Hermansky-Pudlak syndrome 4 (HPS4), transcript variant 3, r
 NM_152841 Homo sapiens Hermansky-Pudlak syndrome 4 (HPS4), transcript variant 2, r
 NM_152842 Homo sapiens Hermansky-Pudlak syndrome 4 (HPS4), transcript variant 5, r
 NM_152843 Homo sapiens Hermansky-Pudlak syndrome 4 (HPS4), transcript variant 4, r
 NM_152850 Homo sapiens phosphatidylinositol glycan, class O (PIGO), transcript variant
 NM_152851 Homo sapiens membrane-spanning 4-domains, subfamily A, member 6A (M
 NM_152852 Homo sapiens membrane-spanning 4-domains, subfamily A, member 6A (M
 NM_152854 Homo sapiens tumor necrosis factor receptor superfamily, member 5 (TNFR
 NM_152855 Homo sapiens immunoglobulin lambda-like polypeptide 1 (IGLL1), transcript
 NM_152856 Homo sapiens RNA binding motif protein 10 (RBM10), transcript variant 2, m
 NM_152857 Homo sapiens Wilms tumor 1 associated protein (WTAP), transcript variant 2
 NM_152858 Homo sapiens Wilms tumor 1 associated protein (WTAP), transcript variant 1
 NM_152860 Homo sapiens Sp7 transcription factor (SP7), mRNA
 NM_152862 Homo sapiens actin related protein 2/3 complex, subunit 2, 34kDa (ARPC2),
 NM_152864 Homo sapiens chromosome 20 open reading frame 58 (C20orf58), mRNA
 NM_152866 Homo sapiens membrane-spanning 4-domains, subfamily A, member 1 (MS
 NM_152867 Homo sapiens membrane-spanning 4-domains, subfamily A, member 1 (MS
 NM_152868 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 4
 NM_152869 Homo sapiens regucalcin (senescence marker protein-30) (RGN), transcript
 NM_152870 Homo sapiens abhydrolase domain containing 1 (ABHD1), transcript variant
 NM_152871 Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFR
 NM_152872 Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFR
 NM_152873 Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFR
 NM_152874 Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFR
 NM_152875 Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFR
 NM_152876 Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFR
 NM_152877 Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFR
 NM_152878 Homo sapiens v-maf musculoaponeurotic fibrosarcoma oncogene homolog f
 NM_152879 Homo sapiens diacylglycerol kinase, delta 130kDa (DGKD), transcript varian
 NM_152880 Homo sapiens PTK7 protein tyrosine kinase 7 (PTK7), transcript variant PTK
 NM_152881 Homo sapiens PTK7 protein tyrosine kinase 7 (PTK7), transcript variant PTK
 NM_152882 Homo sapiens PTK7 protein tyrosine kinase 7 (PTK7), transcript variant PTK
 NM_152883 Homo sapiens PTK7 protein tyrosine kinase 7 (PTK7), transcript variant PTK
 NM_152888 Homo sapiens collagen, type XXII, alpha 1 (COL22A1), mRNA
 NM_152889 Homo sapiens carbohydrate (chondroitin 4) sulfotransferase 13 (CHST13), m
 NM_152890 Homo sapiens collagen, type XXIV, alpha 1 (COL24A1), mRNA
 NM_152891 Homo sapiens protease, serine, 33 (PRSS33), mRNA
 NM_152892 Homo sapiens hypothetical protein DKFZp434K1815 (DKFZp434K1815), mF
 NM_152896 Homo sapiens ubiquitin-like, containing PHD and RING finger domains, 2 (U
 NM_152897 Homo sapiens chromosome 20 open reading frame 161 (C20orf161), transc
 NM_152898 Homo sapiens Fer3-like (Drosophila) (FERD3L), mRNA
 NM_152899 Homo sapiens interleukin 4 induced 1 (IL4I1), transcript variant 1, mRNA
 NM_152900 Homo sapiens membrane-associated guanylate kinase-related (MAGI-3) (M
 NM_152901 Homo sapiens pyrin-domain containing protein 1 (PYC1), mRNA
 NM_152902 Homo sapiens putative MAPK activating protein (MGC3794), mRNA
 NM_152903 Homo sapiens kelch repeat and BTB (POZ) domain containing 6 (KBTBD6),

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NM_153019 Homo sapiens transmembrane protease, serine 6 (TMPRSS6), mRNA
 NM_153020 Homo sapiens RNA binding motif protein 24 (RBM24), mRNA
 NM_153022 Homo sapiens hypothetical protein FLJ31166 (FLJ31166), mRNA
 NM_153023 Homo sapiens spermatogenesis associated 13 (SPATA13), mRNA
 NM_153024 Homo sapiens seven transmembrane helix receptor (FLJ31393), mRNA
 NM_153025 Homo sapiens hypothetical protein FLJ31606 (FLJ31606), mRNA
 NM_153026 Homo sapiens prickly-like 1 (Drosophila) (PRICKLE1), mRNA
 NM_153027 Homo sapiens hypothetical protein FLJ31659 (FLJ31659), mRNA
 NM_153028 Homo sapiens zinc finger protein 75a (ZNF75A), mRNA
 NM_153029 Homo sapiens Nedd4 binding protein 1 (N4BP1), mRNA
 NM_153031 Homo sapiens hypothetical protein FLJ32063 (FLJ32063), mRNA
 NM_153032 Homo sapiens hypothetical protein FLJ32065 (FLJ32065), mRNA
 NM_153033 Homo sapiens potassium channel tetramerisation domain containing 7 (KCT
 NM_153034 Homo sapiens zinc finger protein 488 (ZNF488), mRNA
 NM_153035 Homo sapiens hypothetical protein FLJ32112 (FLJ32112), mRNA
 NM_153036 Homo sapiens chromosome 6 open reading frame 78 (C6orf78), mRNA
 NM_153038 Homo sapiens hypothetical protein FLJ32447 (FLJ32447), mRNA
 NM_153040 Homo sapiens hypothetical protein FLJ32831 (FLJ32831), mRNA
 NM_153041 Homo sapiens hypothetical protein FLJ32955 (FLJ32955), mRNA
 NM_153043 Homo sapiens hypothetical protein FLJ37078 (FLJ37078), mRNA
 NM_153044 Homo sapiens hypothetical protein FLJ35801 (FLJ35801), mRNA
 NM_153045 Homo sapiens chromosome 9 open reading frame 91 (C9orf91), mRNA
 NM_153046 Homo sapiens tudor domain containing 9 (TDRD9), mRNA
 NM_153047 Homo sapiens FYN oncogene related to SRC, FGR, YES (FYN), transcript v
 NM_153048 Homo sapiens FYN oncogene related to SRC, FGR, YES (FYN), transcript v
 NM_153050 Homo sapiens myotubularin related protein 3 (MTMR3), transcript variant 1, i
 NM_153051 Homo sapiens myotubularin related protein 3 (MTMR3), transcript variant 2, i
 NM_153181 Homo sapiens neuropilin (NRP) and tolloid (TLL)-like 1 (NETO1), transcript v
 NM_153182 Homo sapiens MYC induced nuclear antigen (MINA), transcript variant 3, mF
 NM_153183 Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 10
 NM_153184 Homo sapiens immunoglobulin superfamily, member 4D (IGSF4D), mRNA
 NM_153186 Homo sapiens ankyrin repeat domain 15 (ANKRD15), transcript variant 2, mI
 NM_153187 Homo sapiens solute carrier family 22 (organic cation transporter), member 1
 NM_153188 Homo sapiens transportin 1 (TNPO1), transcript variant 2, mRNA
 NM_153189 Homo sapiens sperm adhesion molecule 1 (PH-20 hyaluronidase, zona pellu
 NM_153191 Homo sapiens solute carrier family 22 (organic cation transporter), member 2
 NM_153200 Homo sapiens endothelial differentiation-related factor 1 (EDF1), transcript v
 NM_153201 Homo sapiens heat shock 70kDa protein 8 (HSPA8), transcript variant 2, mR
 NM_153202 Homo sapiens a disintegrin and metalloproteinase domain 33 (ADAM33), tra
 NM_153204 Homo sapiens chromosome 21 open reading frame 90 (C21orf90), mRNA
 NM_153206 Homo sapiens adhesion molecule AMICA (AMICA), mRNA
 NM_153207 Homo sapiens AE binding protein 2 (AEBP2), mRNA
 NM_153208 Homo sapiens hypothetical protein MGC35048 (MGC35048), mRNA
 NM_153209 Homo sapiens hypothetical protein FLJ37300 (FLJ37300), mRNA
 NM_153211 Homo sapiens chromosome 18 open reading frame 17 (C18orf17), mRNA
 NM_153212 Homo sapiens gap junction protein, beta 4 (connexin 30.3) (GJB4), mRNA
 NM_153213 Homo sapiens Rho guanine nucleotide exchange factor (GEF) 19 (ARHGEF
 NM_153214 Homo sapiens hypothetical protein FLJ37440 (FLJ37440), mRNA
 NM_153215 Homo sapiens hypothetical protein FLJ38608 (FLJ38608), mRNA
 NM_153216 Homo sapiens hypothetical protein FLJ25680 (FLJ25680), mRNA
 NM_153217 Homo sapiens hypothetical protein MGC13034 (MGC13034), mRNA
 NM_153218 Homo sapiens hypothetical protein FLJ38725 (FLJ38725), mRNA
 NM_153219 Homo sapiens zinc finger protein 524 (ZNF524), mRNA
 NM_153220 Homo sapiens hypothetical protein MGC35440 (MGC35440), mRNA
 NM_153221 Homo sapiens cartilage intermediate layer protein 2 (CILP2), mRNA
 NM_153223 Homo sapiens hypothetical protein FLJ36090 (FLJ36090), mRNA
 NM_153225 Homo sapiens RPE-spondin (RPESP), mRNA

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NM_153226 Homo sapiens transmembrane protein 20 (TMEM20), mRNA
 NM_153228 Homo sapiens hypothetical protein FLJ38335 (FLJ38335), mRNA
 NM_153229 Homo sapiens hypothetical protein FLJ33318 (FLJ33318), mRNA
 NM_153230 Homo sapiens F-box protein 39 (FBXO39), mRNA
 NM_153231 Homo sapiens zinc finger protein 550 (ZNF550), mRNA
 NM_153232 Homo sapiens CREBBP/EP300 inhibitor 2 (CRI2), mRNA
 NM_153233 Homo sapiens hypothetical protein FLJ36445 (FLJ36445), mRNA
 NM_153234 Homo sapiens chromosome 5 open reading frame 11 (C5orf11), mRNA
 NM_153236 Homo sapiens immune associated nucleotide (hIAN7), mRNA
 NM_153238 Homo sapiens hypothetical protein MGC22001 (MGC22001), mRNA
 NM_153239 Homo sapiens hypothetical protein KIAA1924 (KIAA1924), mRNA
 NM_153240 Homo sapiens nephronophthisis 3 (adolescent) (NPHP3), mRNA
 NM_153244 Homo sapiens chromosome 10 open reading frame 111 (C10orf111), mRNA
 NM_153246 Homo sapiens hypothetical protein MGC45491 (MGC45491), mRNA
 NM_153247 Homo sapiens solute carrier family 29 (nucleoside transporters), member 4 (SLC29A4), mRNA
 NM_153248 Homo sapiens hypothetical protein MGC14276 (MGC14276), mRNA
 NM_153251 Homo sapiens hypothetical protein FLJ25952 (FLJ25952), mRNA
 NM_153252 Homo sapiens bromo domain-containing protein disrupted in leukemia (BROD1), mRNA
 NM_153253 Homo sapiens signal-induced proliferation-associated gene 1 (SIPA1), transcript variant 1, mRNA
 NM_153254 Homo sapiens hypothetical protein FLJ36119 (FLJ36119), mRNA
 NM_153255 Homo sapiens minichromosome maintenance deficient domain containing 1 (MCMDC1), mRNA
 NM_153256 Homo sapiens chromosome 10 open reading frame 47 (C10orf47), mRNA
 NM_153257 Homo sapiens gonadotropin inducible transcription repressor 1 (GIOT-1), mRNA
 NM_153260 Homo sapiens hypothetical protein FLJ36812 (FLJ36812), mRNA
 NM_153261 Homo sapiens hypothetical protein FLJ38101 (FLJ38101), mRNA
 NM_153262 Homo sapiens synaptotagmin XIV (SYT14), mRNA
 NM_153263 Homo sapiens zinc finger protein 549 (ZNF549), mRNA
 NM_153264 Homo sapiens hypothetical protein FLJ35880 (FLJ35880), mRNA
 NM_153265 Homo sapiens hypothetical protein FLJ35827 (FLJ35827), mRNA
 NM_153268 Homo sapiens hypothetical protein MGC33486 (MGC33486), mRNA
 NM_153267 Homo sapiens MAM domain containing 2 (MAMDC2), mRNA
 NM_153268 Homo sapiens hypothetical protein FLJ31579 (FLJ31579), mRNA
 NM_153269 Homo sapiens chromosome 20 open reading frame 96 (C20orf96), mRNA
 NM_153270 Homo sapiens hypothetical protein FLJ34960 (FLJ34960), mRNA
 NM_153271 Homo sapiens hypothetical protein MGC32065 (MGC32065), mRNA
 NM_153273 Homo sapiens inositol hexaphosphate kinase 1 (IHPK1), mRNA
 NM_153274 Homo sapiens vitelliform macular dystrophy 2-like 2 (VMD2L2), mRNA
 NM_153276 Homo sapiens solute carrier family 22 (organic anion transporter), member 6 (SLC22A6), mRNA
 NM_153277 Homo sapiens solute carrier family 22 (organic anion transporter), member 6 (SLC22A6), mRNA
 NM_153278 Homo sapiens solute carrier family 22 (organic anion transporter), member 6 (SLC22A6), mRNA
 NM_153279 Homo sapiens solute carrier family 22 (organic anion transporter), member 6 (SLC22A6), mRNA
 NM_153280 Homo sapiens ubiquitin-activating enzyme E1 (A1S9T and BN75) (UBA1), mRNA
 NM_153281 Homo sapiens hyaluronoglucosaminidase 1 (HYAL1), transcript variant 8, mRNA
 NM_153282 Homo sapiens hyaluronoglucosaminidase 1 (HYAL1), transcript variant 2, mRNA
 NM_153283 Homo sapiens hyaluronoglucosaminidase 1 (HYAL1), transcript variant 3, mRNA
 NM_153284 Homo sapiens hyaluronoglucosaminidase 1 (HYAL1), transcript variant 4, mRNA
 NM_153285 Homo sapiens hyaluronoglucosaminidase 1 (HYAL1), transcript variant 5, mRNA
 NM_153286 Homo sapiens hyaluronoglucosaminidase 1 (HYAL1), transcript variant 6, mRNA
 NM_153289 Homo sapiens defensin, beta 119 (DEFB119), transcript variant 1, mRNA
 NM_153290 Homo sapiens family with sequence similarity 10, member A4 (FAM10A4), mRNA
 NM_153291 Homo sapiens family with sequence similarity 10, member A5 (FAM10A5), mRNA
 NM_153292 Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), transcript variant 1, mRNA
 NM_153320 Homo sapiens solute carrier family 22 (organic anion transporter), member 7 (SLC22A7), mRNA
 NM_153321 Homo sapiens peripheral myelin protein 22 (PMP22), transcript variant 2, mRNA
 NM_153322 Homo sapiens peripheral myelin protein 22 (PMP22), transcript variant 3, mRNA
 NM_153324 Homo sapiens defensin, beta 123 (DEFB123), mRNA
 NM_153325 Homo sapiens defensin, beta 125 (DEFB125), mRNA

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NM_153326 Homo sapiens aldo-keto reductase family 1, member A1 (aldehyde reductase)
 NM_153328 Homo sapiens retinoblastoma binding protein 9 (RBBP9), transcript variant 2
 NM_153329 Homo sapiens hypothetical protein MGC10204 (MGC10204), mRNA
 NM_153330 Homo sapiens DnaJ (Hsp40) homolog, subfamily B, member 8 (DNAJB8), mRNA
 NM_153331 Homo sapiens potassium channel tetramerisation domain containing 6 (KCTD13), mRNA
 NM_153332 Homo sapiens 3' exoribonuclease (3'EXO), mRNA
 NM_153333 Homo sapiens hypothetical protein MGC45400 (MGC45400), mRNA
 NM_153334 Homo sapiens scavenger receptor class F, member 2 (SCARF2), transcript variant 1
 NM_153335 Homo sapiens protein kinase LYK5 (LYK5), mRNA
 NM_153336 Homo sapiens chromosome 10 open reading frame 89 (C10orf89), mRNA
 NM_153337 Homo sapiens selectin ligand interactor cytoplasmic-1 (SLIC1), mRNA
 NM_153338 Homo sapiens hypothetical protein FLJ90165 (FLJ90165), mRNA
 NM_153339 Homo sapiens hypothetical protein FLJ90811 (FLJ90811), mRNA
 NM_153340 Homo sapiens hypothetical protein MGC46534 (MGC46534), mRNA
 NM_153341 Homo sapiens IBR domain containing 3 (IBRDC3), mRNA
 NM_153342 Homo sapiens fasting-inducible integral membrane protein TM6P1 (FLJ9002), mRNA
 NM_153343 Homo sapiens ectonucleotide pyrophosphatase/phosphodiesterase 6 (ENPP6), mRNA
 NM_153344 Homo sapiens chromosome 6 open reading frame 141 (C6orf141), mRNA
 NM_153345 Homo sapiens hypothetical protein FLJ90586 (FLJ90586), mRNA
 NM_153346 Homo sapiens chromosome X open reading frame 20 (CXorf20), mRNA
 NM_153347 Homo sapiens hypothetical protein FLJ90119 (FLJ90119), mRNA
 NM_153348 Homo sapiens F-box and WD-40 domain protein 8 (FBXW8), transcript variant 1
 NM_153350 Homo sapiens F-box and leucine-rich repeat protein 16 (FBXL16), mRNA
 NM_153353 Homo sapiens hypothetical protein MGC27085 (MGC27085), mRNA
 NM_153354 Homo sapiens hypothetical protein MGC33214 (MGC33214), mRNA
 NM_153355 Homo sapiens T-cell lymphoma breakpoint associated target 1 (TCBA1), mRNA
 NM_153357 Homo sapiens hypothetical protein MGC34741 (MGC34741), mRNA
 NM_153358 Homo sapiens solute carrier family 16 (monocarboxylic acid transporters), member 1
 NM_153359 Homo sapiens hypothetical protein FLJ90396 (FLJ90396), mRNA
 NM_153360 Homo sapiens hypothetical protein MGC24975 (MGC24975), mRNA
 NM_153361 Homo sapiens hypothetical protein FLJ90166 (FLJ90166), mRNA
 NM_153362 Homo sapiens hypothetical protein MGC42105 (MGC42105), mRNA
 NM_153363 Homo sapiens protease, serine, 35 (PRSS35), mRNA
 NM_153364 Homo sapiens hypothetical protein MGC42415 (MGC42415), mRNA
 NM_153365 Homo sapiens hypothetical protein MGC39520 (MGC39520), mRNA
 NM_153366 Homo sapiens hypothetical protein FLJ90013 (FLJ90013), mRNA
 NM_153367 Homo sapiens chromosome 10 open reading frame 56 (C10orf56), mRNA
 NM_153368 Homo sapiens connexin40.1 (CX40.1), mRNA
 NM_153369 Homo sapiens KIAA1919 (KIAA1919), mRNA
 NM_153370 Homo sapiens protease inhibitor 16 (PI16), mRNA
 NM_153371 Homo sapiens ligand of numb-protein X 2 (LNX2), mRNA
 NM_153373 Homo sapiens hypothetical protein MGC15875 (MGC15875), mRNA
 NM_153374 Homo sapiens hypothetical protein MGC35274 (MGC35274), mRNA
 NM_153375 Homo sapiens placenta-specific 2 (PLAC2), mRNA
 NM_153376 Homo sapiens hypothetical protein FLJ90575 (FLJ90575), mRNA
 NM_153377 Homo sapiens leucine-rich repeats and immunoglobulin-like domains 3 (LRRC13), mRNA
 NM_153378 Homo sapiens solute carrier family 22 (organic anion/cation transporter), member 1
 NM_153379 Homo sapiens kringle containing transmembrane protein 1 (KREMEN1), mRNA
 NM_153380 Homo sapiens zinc finger protein 41 (ZNF41), transcript variant 2, mRNA
 NM_153381 Homo sapiens pro-melanin-concentrating hormone-like 2 (PMCHL2), mRNA
 NM_153425 Homo sapiens TNFRSF1A-associated via death domain (TRADD), transcript variant 1
 NM_153426 Homo sapiens paired-like homeodomain transcription factor 2 (PITX2), transcript variant 1
 NM_153427 Homo sapiens paired-like homeodomain transcription factor 2 (PITX2), transcript variant 2
 NM_153437 Homo sapiens outer dense fiber of sperm tails 2 (ODF2), transcript variant 2, mRNA
 NM_153442 Homo sapiens G protein-coupled receptor 26 (GPR26), mRNA
 NM_153443 Homo sapiens killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail
 NM_153444 Homo sapiens olfactory receptor, family 5, subfamily P, member 2 (OR5P2), mRNA

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NM_153445 Homo sapiens olfactory receptor, family 5, subfamily P, member 3 (OR5P3),
 NM_153446 Homo sapiens UDP-GalNAc:Neu5Acalpha2-3Galbeta-R beta1,4-N-acetylgal
 NM_153447 Homo sapiens NACHT, leucine rich repeat and PYD containing 5 (NALP5), n
 NM_153448 Homo sapiens extraembryonic, spermatogenesis, homeobox 1-like (ESX1L),
 NM_153449 Homo sapiens solute carrier family 2 (facilitated glucose transporter), membe
 NM_153450 Homo sapiens lung cancer metastasis-related protein 1 (LCMR1), mRNA
 NM_153451 Homo sapiens oral cancer overexpressed 1 (ORAOV1), mRNA
 NM_153453 Homo sapiens vestigial like 2 (Drosophila) (VGLL2), transcript variant 2, mRf
 NM_153454 Homo sapiens chromosome 21 open reading frame 86 (C21orf86), mRNA
 NM_153456 Homo sapiens heparan sulfate 6-O-sulfotransferase 3 (HS6ST3), mRNA
 NM_153460 Homo sapiens interleukin 17 receptor C (IL17RC), transcript variant 2, mRN/
 NM_153461 Homo sapiens interleukin 17 receptor C (IL17RC), transcript variant 1, mRN/
 NM_153462 Homo sapiens interleukin 17 receptor C (IL17RC), transcript variant 4, mRN/
 NM_153463 Homo sapiens interleukin 17 receptor C (IL17RC), transcript variant 5, mRN/
 NM_153464 Homo sapiens interleukin enhancer binding factor 3, 90kDa (ILF3), transcript
 NM_153477 Homo sapiens ubiquitously-expressed transcript (UXT), transcript variant 1, r
 NM_153478 Homo sapiens chondrosarcoma associated gene 1 (CSAG1), transcript varia
 NM_153479 Homo sapiens chondrosarcoma associated gene 1 (CSAG1), transcript varia
 NM_153480 Homo sapiens interleukin 17 receptor E (IL17RE), transcript variant 1, mRN/
 NM_153481 Homo sapiens interleukin 17 receptor E (IL17RE), transcript variant 2, mRN/
 NM_153482 Homo sapiens interleukin 17 receptor E (IL17RE), transcript variant 4, mRN/
 NM_153483 Homo sapiens interleukin 17 receptor E (IL17RE), transcript variant 5, mRN/
 NM_153485 Homo sapiens nucleoporin 155kDa (NUP155), transcript variant 1, mRNA
 NM_153486 Homo sapiens lactate dehydrogenase D (LDHD), nuclear gene encoding mit
 NM_153487 Homo sapiens MAM domain containing glycosylphosphatidylinositol anchor 1
 NM_153488 Homo sapiens melanoma antigen, family A, 2B (MAGEA2B), mRNA
 NM_153490 Homo sapiens keratin 13 (KRT13), transcript variant 1, mRNA
 NM_153497 Homo sapiens mitogen-activated protein kinase kinase kinase 7 interacting p
 NM_153498 Homo sapiens calcium/calmodulin-dependent protein kinase ID (CAMK1D), t
 NM_153499 Homo sapiens calcium/calmodulin-dependent protein kinase kinase 2, beta
 NM_153500 Homo sapiens calcium/calmodulin-dependent protein kinase kinase 2, beta
 NM_153503 Homo sapiens component of oligomeric golgi complex 7 (COG7), mRNA
 NM_153604 Homo sapiens myocardin (MYOCD), mRNA
 NM_153606 Homo sapiens hypothetical protein FLJ32796 (FLJ32796), mRNA
 NM_153607 Homo sapiens adult retina protein (LOC153222), mRNA
 NM_153608 Homo sapiens hypothetical protein MGC17986 (MGC17986), mRNA
 NM_153609 Homo sapiens transmembrane protease, serine 6 (TMPRSS6), mRNA
 NM_153610 Homo sapiens cardiomyopathy associated 5 (CMYA5), mRNA
 NM_153611 Homo sapiens hypothetical protein MGC20446 (MGC20446), mRNA
 NM_153612 Homo sapiens heparan sulfate (glucosamine) 3-O-sulfotransferase 5 (HS3S1
 NM_153613 Homo sapiens PISC domain containing hypothetical protein (LOC254531), r
 NM_153614 Homo sapiens testis spermatogenesis apoptosis-related protein 6 (TSARG6)
 NM_153615 Homo sapiens Ral-GDS related protein Rgr (Rgr), mRNA
 NM_153616 Homo sapiens sema domain, transmembrane domain (TM), and cytoplasmic
 NM_153617 Homo sapiens sema domain, transmembrane domain (TM), and cytoplasmic
 NM_153618 Homo sapiens sema domain, transmembrane domain (TM), and cytoplasmic
 NM_153619 Homo sapiens sema domain, transmembrane domain (TM), and cytoplasmic
 NM_153620 Homo sapiens homeo box A1 (HOXA1), transcript variant 2, mRNA
 NM_153631 Homo sapiens homeo box A3 (HOXA3), transcript variant 2, mRNA
 NM_153632 Homo sapiens homeo box A3 (HOXA3), transcript variant 3, mRNA
 NM_153633 Homo sapiens homeo box C4 (HOXC4), transcript variant 2, mRNA
 NM_153634 Homo sapiens copine VIII (CPNE8), mRNA
 NM_153635 Homo sapiens copine family member (LOC151835), mRNA
 NM_153636 Homo sapiens copine VII (CPNE7), transcript variant 1, mRNA
 NM_153637 Homo sapiens pantothenate kinase 2 (Hallervorden-Spatz syndrome) (PANK
 NM_153638 Homo sapiens pantothenate kinase 2 (Hallervorden-Spatz syndrome) (PANK
 NM_153639 Homo sapiens pantothenate kinase 2 (Hallervorden-Spatz syndrome) (PANK

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NM_153640 Homo sapiens pantothenate kinase 2 (Hallervorden-Spatz syndrome) (PANK)
 NM_153641 Homo sapiens pantothenate kinase 2 (Hallervorden-Spatz syndrome) (PANK)
 NM_153645 Homo sapiens nucleoporin 50kDa (NUP50), transcript variant 3, mRNA
 NM_153646 Homo sapiens solute carrier family 24 (sodium/potassium/calcium exchanger)
 NM_153647 Homo sapiens solute carrier family 24 (sodium/potassium/calcium exchanger)
 NM_153648 Homo sapiens solute carrier family 24 (sodium/potassium/calcium exchanger)
 NM_153649 Homo sapiens tropomyosin 3 (TPM3), mRNA
 NM_153675 Homo sapiens forkhead box A2 (FOXA2), transcript variant 2, mRNA
 NM_153676 Homo sapiens Usher syndrome critical region 1C (autosomal recessive, severe) (USH1C),
 NM_153681 Homo sapiens Down syndrome critical region gene 5 (DSCR5), transcript var
 NM_153682 Homo sapiens Down syndrome critical region gene 5 (DSCR5), transcript var
 NM_153683 Homo sapiens klotho (KL), transcript variant 2, mRNA
 NM_153684 Homo sapiens nucleoporin 50kDa (NUP50), transcript variant 1, mRNA
 NM_153685 Homo sapiens hypothetical protein DKFZp547D2210 (DKFZp547D2210), mF
 NM_153686 Homo sapiens transcription factor MLR1 (MLR1), mRNA
 NM_153687 Homo sapiens hypothetical protein FLJ31051 (FLJ31051), mRNA
 NM_153688 Homo sapiens zinc finger protein 1 homolog (mouse) (ZFP1), mRNA
 NM_153689 Homo sapiens hypothetical protein FLJ38973 (FLJ38973), mRNA
 NM_153690 Homo sapiens family with sequence similarity 43, member A (FAM43A), mRf
 NM_153691 Homo sapiens hypothetical protein FLJ90036 (FLJ90036), mRNA
 NM_153692 Homo sapiens hypothetical protein FLJ90724 (FLJ90724), mRNA
 NM_153693 Homo sapiens homeo box C6 (HOXC6), transcript variant 2, mRNA
 NM_153694 Homo sapiens synaptonemal complex protein 3 (SYCP3), mRNA
 NM_153695 Homo sapiens zinc finger protein 367 (ZNF367), mRNA
 NM_153696 Homo sapiens prostate-specific membrane antigen-like protein (PSMAL/GCF
 NM_153697 Homo sapiens hypothetical protein DKFZp434D2328 (LOC91526), mRNA
 NM_153699 Homo sapiens glutathione S-transferase A5 (GSTA5), mRNA
 NM_153700 Homo sapiens stereocilin (STRC), mRNA
 NM_153701 Homo sapiens interleukin 12 receptor, beta 1 (IL12RB1), transcript variant 2,
 NM_153702 Homo sapiens hypothetical protein MGC10084 (MGC10084), mRNA
 NM_153703 Homo sapiens podocan (PODN), mRNA
 NM_153704 Homo sapiens hypothetical protein MGC26979 (MGC26979), mRNA
 NM_153705 Homo sapiens KDEL (Lys-Asp-Glu-Leu) containing 2 (KDEL2), mRNA
 NM_153706 Homo sapiens hypothetical protein MGC33648 (MGC33648), mRNA
 NM_153707 Homo sapiens chromosome 9 open reading frame 138 (C9orf138), mRNA
 NM_153708 Homo sapiens hypothetical protein MGC35450 (MGC35450), mRNA
 NM_153709 Homo sapiens hypothetical protein MGC40168 (MGC40168), mRNA
 NM_153711 Homo sapiens chromosome 6 open reading frame 188 (C6orf188), mRNA
 NM_153712 Homo sapiens tubulin tyrosine ligase (TTL), mRNA
 NM_153713 Homo sapiens hypothetical protein MGC46719 (MGC46719), mRNA
 NM_153714 Homo sapiens chromosome 10 open reading frame 67 (C10orf67), mRNA
 NM_153715 Homo sapiens homeo box A10 (HOXA10), transcript variant 2, mRNA
 NM_153716 Homo sapiens heat shock transcription factor, Y linked 2 (HSFY2), transcript
 NM_153717 Homo sapiens Ellis van Creveld syndrome (EVC), transcript variant 2, mRNA
 NM_153718 Homo sapiens nucleoporin 62kDa (NUP62), transcript variant 3, mRNA
 NM_153719 Homo sapiens nucleoporin 62kDa (NUP62), transcript variant 1, mRNA
 NM_153741 Homo sapiens dolichyl-phosphate mannosyltransferase polypeptide 3 (DPMK
 NM_153742 Homo sapiens cystathionase (cystathionine gamma-lyase) (CTH), transcript
 NM_153746 Homo sapiens zinc finger, DHHC domain containing 14 (ZDHHC14), mRNA
 NM_153747 Homo sapiens phosphatidylinositol glycan, class C (PIGC), transcript variant
 NM_153748 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, m
 NM_153750 Homo sapiens chromosome 21 open reading frame 81 (C21orf81), mRNA
 NM_153752 Homo sapiens chromosome 21 open reading frame 84 (C21orf84), mRNA
 NM_153754 Homo sapiens chromosome 21 open reading frame 88 (C21orf88), mRNA
 NM_153756 Homo sapiens fibronectin type III domain containing 5 (FNDC5), mRNA
 NM_153757 Homo sapiens nucleosome assembly protein 1-like 5 (NAP1L5), mRNA
 NM_153758 Homo sapiens interleukin 19 (IL19), transcript variant 1, mRNA

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NM_153759 Homo sapiens DNA (cytosine-5)-methyltransferase 3 alpha (DNMT3A), trans
 NM_153763 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, mu
 NM_153764 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 1
 NM_153765 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 1
 NM_153766 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 1
 NM_153767 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 1
 NM_153768 Homo sapiens calcium-binding tyrosine-(Y)-phosphorylation regulated (fibro
 NM_153769 Homo sapiens calcium-binding tyrosine-(Y)-phosphorylation regulated (fibro
 NM_153770 Homo sapiens calcium-binding tyrosine-(Y)-phosphorylation regulated (fibro
 NM_153773 Homo sapiens chromosome 21 open reading frame 99 (C21orf99), mRNA
 NM_153809 Homo sapiens TAF1-like RNA polymerase II, TATA box binding protein (TBP
 NM_153810 Homo sapiens chromosome 10 open reading frame 46 (C10orf46), mRNA
 NM_153811 Homo sapiens solute carrier family 38, member 6 (SLC38A6), mRNA
 NM_153812 Homo sapiens PHD finger protein 13 (PHF13), mRNA
 NM_153813 Homo sapiens zinc finger protein, multitype 1 (ZFPM1), mRNA
 NM_153815 Homo sapiens Ras protein-specific guanine nucleotide-releasing factor 1 (R/
 NM_153816 Homo sapiens sorting nexin 14 (SNX14), transcript variant 1, mRNA
 NM_153818 Homo sapiens peroxisome biogenesis factor 10 (PEX10), transcript variant 1
 NM_153819 Homo sapiens RAS guanyl releasing protein 2 (calcium and DAG-regulated)
 NM_153822 Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase,
 NM_153823 Homo sapiens germ cell associated 1 (GSG1), mRNA
 NM_153824 Homo sapiens pyrroline-5-carboxylate reductase 1 (PYCR1), transcript varia
 NM_153825 Homo sapiens soluble liver antigen/liver pancreas antigen (SLA/LP), mRNA
 NM_153826 Homo sapiens membrane cofactor protein (CD46, trophoblast-lymphocyte co
 NM_153827 Homo sapiens misshapen/NIK-related kinase (MINK), transcript variant 3, m
 NM_153828 Homo sapiens reticulin 4 (RTN4), transcript variant 2, mRNA
 NM_153831 Homo sapiens PTK2 protein tyrosine kinase 2 (PTK2), transcript variant 1, m
 NM_153832 Homo sapiens G protein-coupled receptor 161 (GPR161), mRNA
 NM_153833 Homo sapiens H1 histone family, member O, oocyte-specific (H1FOO), mRN
 NM_153834 Homo sapiens G protein-coupled receptor 112 (GPR112), mRNA
 NM_153835 Homo sapiens G protein-coupled receptor 113 (GPR113), mRNA
 NM_153836 Homo sapiens cellular repressor of E1A-stimulated genes 2 (CREG2), mRN/
 NM_153837 Homo sapiens G protein-coupled receptor 114 (GPR114), mRNA
 NM_153838 Homo sapiens G protein-coupled receptor 115 (GPR115), mRNA
 NM_153839 Homo sapiens G protein-coupled receptor 111 (GPR111), mRNA
 NM_153840 Homo sapiens G protein-coupled receptor 110 (GPR110), mRNA
 NM_156036 Homo sapiens homeo box B6 (HOXB6), transcript variant 3, mRNA
 NM_156037 Homo sapiens homeo box B6 (HOXB6), transcript variant 1, mRNA
 NM_156038 Homo sapiens colony stimulating factor 3 receptor (granulocyte) (CSF3R), tr
 NM_156039 Homo sapiens colony stimulating factor 3 receptor (granulocyte) (CSF3R), tr
 NM_170587 Homo sapiens regulator of G-protein signalling 20 (RGS20), mRNA
 NM_170589 Homo sapiens AF15q14 protein (AF15Q14), mRNA
 NM_170600 Homo sapiens SH2 domain containing 3C (SH2D3C), mRNA
 NM_170601 Homo sapiens cytosolic sialic acid 9-O-acetyltransferase homolog (CSE-C), mF
 NM_170602 Homo sapiens RAS guanyl releasing protein 4 (RASGRP4), transcript varian
 NM_170603 Homo sapiens RAS guanyl releasing protein 4 (RASGRP4), transcript varian
 NM_170604 Homo sapiens RAS guanyl releasing protein 4 (RASGRP4), transcript varian
 NM_170605 Homo sapiens InaD-like protein (INADL), transcript variant 1, mRNA
 NM_170606 Homo sapiens myeloid/lymphoid or mixed-lineage leukemia 3 (MLL3), mRNA
 NM_170607 Homo sapiens transcription factor-like 4 (TCFL4), transcript variant 3, mRNA
 NM_170609 Homo sapiens cysteine-rich secretory protein 1 (CRISP1), transcript variant
 NM_170610 Homo sapiens histone 1, H2ba (HIST1H2BA), mRNA
 NM_170662 Homo sapiens Cas-BR-M (murine) ecotropic retroviral transforming sequence
 NM_170663 Homo sapiens misshapen/NIK-related kinase (MINK), transcript variant 2, m
 NM_170664 Homo sapiens otocorin (OTOA), mRNA
 NM_170665 Homo sapiens ATPase, Ca++ transporting, cardiac muscle, slow twitch 2 (AT
 NM_170672 Homo sapiens RAS guanyl releasing protein 3 (calcium and DAG-regulated)

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NM_170674 Homo sapiens Meis1, myeloid ecotropic viral integration site 1 homolog 2 (m
 NM_170675 Homo sapiens Meis1, myeloid ecotropic viral integration site 1 homolog 2 (m
 NM_170676 Homo sapiens Meis1, myeloid ecotropic viral integration site 1 homolog 2 (m
 NM_170677 Homo sapiens Meis1, myeloid ecotropic viral integration site 1 homolog 2 (m
 NM_170678 Homo sapiens integrin beta 1 binding protein 3 (ITGB1BP3), mRNA
 NM_170679 Homo sapiens S-phase kinase-associated protein 1A (p19A) (SKP1A), transcr
 NM_170681 Homo sapiens mitochondrial elongation factor G2 (EFG2), nuclear gene enc
 NM_170682 Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 2 (P2RX2)
 NM_170683 Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 2 (P2RX2)
 NM_170685 Homo sapiens tachykinin 4 (hemokinin) (TAC4), mRNA
 NM_170686 Homo sapiens zinc finger protein 398 (ZNF398), transcript variant 1, mRNA
 NM_170691 Homo sapiens mitochondrial elongation factor G2 (EFG2), nuclear gene enc
 NM_170692 Homo sapiens RAS protein activator like 2 (RASAL2), transcript variant 2, m
 NM_170693 Homo sapiens serum/glucocorticoid regulated kinase 2 (SGK2), transcript va
 NM_170694 Homo sapiens serine hydrolase-like (SERHL), mRNA
 NM_170695 Homo sapiens TGFB-Induced factor (TALE family homeobox) (TGIF), transcr
 NM_170696 Homo sapiens aldehyde dehydrogenase 1 family, member A2 (ALDH1A2), tr
 NM_170697 Homo sapiens aldehyde dehydrogenase 1 family, member A2 (ALDH1A2), tr
 NM_170698 Homo sapiens similar to CGI-96 (dJ22E13.2), mRNA
 NM_170699 Homo sapiens G protein-coupled bile acid receptor 1 (GPBAR1), mRNA
 NM_170705 Homo sapiens isoprenylcysteine carboxyl methyltransferase (ICMT), transcrip
 NM_170706 Homo sapiens nicotinamide nucleotide adenyltransferase 2 (NMNAT2), tra
 NM_170707 Homo sapiens lamin A/C (LMNA), transcript variant 1, mRNA
 NM_170708 Homo sapiens lamin A/C (LMNA), transcript variant 3, mRNA
 NM_170709 Homo sapiens serum/glucocorticoid regulated kinase-like (SGKL), transcript
 NM_170710 Homo sapiens WD repeat domain 17 (WDR17), transcript variant 1, mRNA
 NM_170711 Homo sapiens DAZ associated protein 1 (DAZAP1), transcript variant 1, mR
 NM_170712 Homo sapiens Ras association (RalGDS/AF-6) domain family 1 (RASSF1), t
 NM_170713 Homo sapiens Ras association (RalGDS/AF-6) domain family 1 (RASSF1), t
 NM_170714 Homo sapiens Ras association (RalGDS/AF-6) domain family 1 (RASSF1), t
 NM_170715 Homo sapiens Ras association (RalGDS/AF-6) domain family 1 (RASSF1), t
 NM_170716 Homo sapiens Ras association (RalGDS/AF-6) domain family 1 (RASSF1), t
 NM_170717 Homo sapiens Ras association (RalGDS/AF-6) domain family 1 (RASSF1), t
 NM_170719 Homo sapiens chromosome 13 open reading frame 23 (C13orf23), transcript
 NM_170720 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 1
 NM_170721 Homo sapiens musashi homolog 2 (Drosophila) (MSI2), transcript variant 2, i
 NM_170722 Homo sapiens NOD9 protein (NOD9), transcript variant 2, mRNA
 NM_170723 Homo sapiens chromodomain protein, Y-linked, 1 (CDY1), transcript variant
 NM_170724 Homo sapiens polycystic kidney and hepatic disease 1 (autosomal recessive
 NM_170725 Homo sapiens piggyBac transposable element derived 2 (PGBD2), mRNA
 NM_170726 Homo sapiens aldehyde dehydrogenase 4 family, member A1 (ALDH4A1), n
 NM_170731 Homo sapiens brain-derived neurotrophic factor (BDNF), transcript variant 3,
 NM_170732 Homo sapiens brain-derived neurotrophic factor (BDNF), transcript variant 2,
 NM_170733 Homo sapiens brain-derived neurotrophic factor (BDNF), transcript variant 5,
 NM_170734 Homo sapiens brain-derived neurotrophic factor (BDNF), transcript variant 6,
 NM_170735 Homo sapiens brain-derived neurotrophic factor (BDNF), transcript variant 1,
 NM_170736 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 1
 NM_170737 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 1
 NM_170738 Homo sapiens mitochondrial ribosomal protein L11 (MRPL11), nuclear gene
 NM_170739 Homo sapiens mitochondrial ribosomal protein L11 (MRPL11), nuclear gene
 NM_170740 Homo sapiens aldehyde dehydrogenase 5 family, member A1 (succinate-ser
 NM_170741 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 1
 NM_170742 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 1
 NM_170743 Homo sapiens interleukin 28 receptor, alpha (interferon, lambda receptor) (IL
 NM_170744 Homo sapiens unc-5 homolog B (C. elegans) (UNC5B), mRNA
 NM_170745 Homo sapiens histone 1, H2aa (HIST1H2AA), mRNA
 NM_170746 Homo sapiens chromosome 11 open reading frame 31 (C11orf31), mRNA

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NM_170750 Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase,
 NM_170751 Homo sapiens chromodomain protein, Y-like (CDYL), transcript variant 2, mf
 NM_170752 Homo sapiens chromodomain protein, Y-like (CDYL), transcript variant 3, mf
 NM_170753 Homo sapiens piggyBac transposable element derived 3 (PGBD3), mRNA
 NM_170754 Homo sapiens tensin like C1 domain containing phosphatase (TENC1), trans
 NM_170768 Homo sapiens zinc finger protein 91 homolog (mouse) (ZFP91), transcript va
 NM_170769 Homo sapiens ring finger protein 39 (RNF39), transcript variant 2, mRNA
 NM_170770 Homo sapiens ring finger protein 39 (RNF39), transcript variant 3, mRNA
 NM_170771 Homo sapiens aldehyde dehydrogenase 8 family, member A1 (ALDH8A1), tr
 NM_170773 Homo sapiens Ras association (RalGDS/AF-6) domain family 2 (RASSF2), b
 NM_170774 Homo sapiens Ras association (RalGDS/AF-6) domain family 2 (RASSF2), b
 NM_170775 Homo sapiens potassium intermediate/small conductance calcium-activated
 NM_170776 Homo sapiens G protein-coupled receptor 97 (GPR97), mRNA
 NM_170780 Homo sapiens MOX2 receptor (MOX2R), transcript variant 4, mRNA
 NM_170781 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 11 (PPP1
 NM_170782 Homo sapiens potassium intermediate/small conductance calcium-activated
 NM_170783 Homo sapiens zinc ribbon domain containing, 1 (ZNRD1), transcript variant 2
 NM_170784 Homo sapiens McKusick-Kaufman syndrome (MKKS), transcript variant 2, m
 NM_171825 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
 NM_171827 Homo sapiens CD8 antigen, alpha polypeptide (p32) (CD8A), transcript varia
 NM_171828 Homo sapiens potassium large conductance calcium-activated channel, subf
 NM_171829 Homo sapiens potassium large conductance calcium-activated channel, subf
 NM_171830 Homo sapiens potassium large conductance calcium-activated channel, subf
 NM_171846 Homo sapiens lactamase, beta (LACTB), nuclear gene encoding mitochondr
 NM_171982 Homo sapiens tripartite motif-containing 35 (TRIM35), transcript variant 2, m
 NM_171997 Homo sapiens ubiquitin specific protease 2 (USP2), transcript variant 2, mR
 NM_171998 Homo sapiens RAB39B, member RAS oncogene family (RAB39B), mRNA
 NM_171999 Homo sapiens sal-like 3 (Drosophila) (SALL3), mRNA
 NM_172000 Homo sapiens putative membrane protein HE9 (HE9), mRNA
 NM_172002 Homo sapiens J-type co-chaperone HSC20 (HSC20), mRNA
 NM_172003 Homo sapiens COBW domain containing 2 (CBWD2), mRNA
 NM_172004 Homo sapiens dendritic cell-associated lectin-1 (DICAL1), mRNA
 NM_172005 Homo sapiens WAP four-disulfide core domain 13 (WFDC13), mRNA
 NM_172006 Homo sapiens WAP four-disulfide core domain 10B (WFDC10B), transcript v
 NM_172014 Homo sapiens tumor necrosis factor (ligand) superfamily, member 14 (TNFS
 NM_172016 Homo sapiens tripartite motif-containing 39 (TRIM39), transcript variant 2, m
 NM_172020 Homo sapiens POM121 membrane glycoprotein (rat) (POM121), mRNA
 NM_172024 Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 13
 NM_172025 Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 13
 NM_172026 Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 13
 NM_172027 Homo sapiens ankyrin repeat and BTB (POZ) domain containing 1 (ABTB1),
 NM_172028 Homo sapiens ankyrin repeat and BTB (POZ) domain containing 1 (ABTB1),
 NM_172037 Homo sapiens retinol dehydrogenase 10 (all-trans) (RDH10), mRNA
 NM_172056 Homo sapiens potassium voltage-gated channel, subfamily H (eag-related), i
 NM_172057 Homo sapiens potassium voltage-gated channel, subfamily H (eag-related), i
 NM_172058 Homo sapiens eyes absent homolog 1 (Drosophila) (EYA1), transcript varian
 NM_172059 Homo sapiens eyes absent homolog 1 (Drosophila) (EYA1), transcript varian
 NM_172060 Homo sapiens eyes absent homolog 1 (Drosophila) (EYA1), transcript varian
 NM_172069 Homo sapiens pleckstrin homology domain containing, family H (with IlyTH4
 NM_172070 Homo sapiens similar to F10G7.10.p (KIAA2024), mRNA
 NM_172078 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
 NM_172079 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
 NM_172080 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
 NM_172081 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
 NM_172082 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
 NM_172083 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
 NM_172084 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II

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NM_172087 Homo sapiens tumor necrosis factor (ligand) superfamily, member 13 (TNFS
 NM_172088 Homo sapiens tumor necrosis factor (ligand) superfamily, member 13 (TNFS
 NM_172089 Homo sapiens tumor necrosis factor (ligand) superfamily, member 12-memb
 NM_172095 Homo sapiens cation channel, sperm associated 2 (CATSPER2), transcript v
 NM_172096 Homo sapiens cation channel, sperm associated 2 (CATSPER2), transcript v
 NM_172097 Homo sapiens cation channel, sperm associated 2 (CATSPER2), transcript v
 NM_172098 Homo sapiens eyes absent homolog 3 (Drosophila) (EYA3), transcript varian
 NM_172099 Homo sapiens CD8 antigen, beta polypeptide 1 (p37) (CD8B1), transcript vai
 NM_172100 Homo sapiens CD8 antigen, beta polypeptide 1 (p37) (CD8B1), transcript vai
 NM_172101 Homo sapiens CD8 antigen, beta polypeptide 1 (p37) (CD8B1), transcript vai
 NM_172102 Homo sapiens CD8 antigen, beta polypeptide 1 (p37) (CD8B1), transcript vai
 NM_172103 Homo sapiens eyes absent homolog 4 (Drosophila) (EYA4), transcript varian
 NM_172104 Homo sapiens eyes absent homolog 4 (Drosophila) (EYA4), transcript varian
 NM_172105 Homo sapiens eyes absent homolog 4 (Drosophila) (EYA4), transcript varian
 NM_172106 Homo sapiens potassium voltage-gated channel, KQT-like subfamily, membe
 NM_172107 Homo sapiens potassium voltage-gated channel, KQT-like subfamily, membe
 NM_172108 Homo sapiens potassium voltage-gated channel, KQT-like subfamily, membe
 NM_172109 Homo sapiens potassium voltage-gated channel, KQT-like subfamily, membe
 NM_172110 Homo sapiens eyes absent homolog 2 (Drosophila) (EYA2), transcript varian
 NM_172111 Homo sapiens eyes absent homolog 2 (Drosophila) (EYA2), transcript varian
 NM_172112 Homo sapiens eyes absent homolog 2 (Drosophila) (EYA2), transcript varian
 NM_172113 Homo sapiens eyes absent homolog 2 (Drosophila) (EYA2), transcript varian
 NM_172115 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
 NM_172127 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
 NM_172128 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
 NM_172130 Homo sapiens potassium voltage-gated channel, shaker-related subfamily, b
 NM_172131 Homo sapiens WAP four-disulfide core domain 10B (WFDC10B), transcript v
 NM_172138 Homo sapiens interleukin 28A (interferon, lambda 2) (IL28A), mRNA
 NM_172139 Homo sapiens interleukin 28B (interferon, lambda 3) (IL28B), mRNA
 NM_172140 Homo sapiens interleukin 29 (interferon, lambda 1) (IL29), mRNA
 NM_172159 Homo sapiens potassium voltage-gated channel, shaker-related subfamily, b
 NM_172160 Homo sapiens potassium voltage-gated channel, shaker-related subfamily, b
 NM_172163 Homo sapiens potassium voltage-gated channel, KQT-like subfamily, membe
 NM_172164 Homo sapiens nuclear autoantigenic sperm protein (histone-binding) (NASP)
 NM_172185 Homo sapiens mutS homolog 5 (E. coli) (MSH5), transcript variant 2, mRNA
 NM_172186 Homo sapiens mutS homolog 5 (E. coli) (MSH5), transcript variant 4, mRNA
 NM_172167 Homo sapiens NADPH oxidase organizer 1 (NOXO1), transcript variant b, ml
 NM_172168 Homo sapiens NADPH oxidase organizer 1 (NOXO1), transcript variant c, ml
 NM_172169 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
 NM_172170 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
 NM_172171 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
 NM_172172 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
 NM_172173 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
 NM_172174 Homo sapiens interleukin 15 (IL15), transcript variant 1, mRNA
 NM_172175 Homo sapiens interleukin 15 (IL15), transcript variant 2, mRNA
 NM_172177 Homo sapiens mitochondrial ribosomal protein L42 (MRPL42), nuclear gene
 NM_172178 Homo sapiens mitochondrial ribosomal protein L42 (MRPL42), nuclear gene
 NM_172193 Homo sapiens kelch domain containing 1 (KLHDC1), mRNA
 NM_172195 Homo sapiens eukaryotic translation initiation factor 2B, subunit 4 delta, 67kD
 NM_172196 Homo sapiens TFIIA-alpha/beta-like factor (ALF), transcript variant 2, mRNA
 NM_172197 Homo sapiens advanced glycosylation end product-specific receptor (AGER)
 NM_172198 Homo sapiens potassium voltage-gated channel, Shal-related subfamily, mei
 NM_172199 Homo sapiens adenylate kinase 2 (AK2), transcript variant AK2C, mRNA
 NM_172200 Homo sapiens interleukin 15 receptor, alpha (IL15RA), transcript variant 2, r
 NM_172201 Homo sapiens potassium voltage-gated channel, Isk-related family, member
 NM_172206 Homo sapiens calcium/calmodulin-dependent protein kinase kinase 1, alpha
 NM_172207 Homo sapiens calcium/calmodulin-dependent protein kinase kinase 1, alpha

NM_172208 Homo sapiens TAP binding protein (tapasin) (TAPBP), transcript variant 2, m
 NM_172209 Homo sapiens TAP binding protein (tapasin) (TAPBP), transcript variant 3, m
 NM_172210 Homo sapiens colony stimulating factor 1 (macrophage) (CSF1), transcript v
 NM_172211 Homo sapiens colony stimulating factor 1 (macrophage) (CSF1), transcript v
 NM_172212 Homo sapiens colony stimulating factor 1 (macrophage) (CSF1), transcript v
 NM_172213 Homo sapiens CD8 antigen, beta polypeptide 1 (p37) (CD8B1), transcript va
 NM_172214 Homo sapiens calcium/calmodulin-dependent protein kinase kinase 2, beta (
 NM_172215 Homo sapiens calcium/calmodulin-dependent protein kinase kinase 2, beta (
 NM_172216 Homo sapiens calcium/calmodulin-dependent protein kinase kinase 2, beta (
 NM_172217 Homo sapiens interleukin 16 (lymphocyte chemoattractant factor) (IL16), tra
 NM_172218 Homo sapiens sperm associated antigen 1 (SPAG1), transcript variant 2, mF
 NM_172219 Homo sapiens colony stimulating factor 3 (granulocyte) (CSF3), transcript va
 NM_172220 Homo sapiens colony stimulating factor 3 (granulocyte) (CSF3), transcript va
 NM_172225 Homo sapiens diencephalon/mesencephalon homeobox 1 (DMBX1), transcri
 NM_172226 Homo sapiens calcium/calmodulin-dependent protein kinase kinase 2, beta (
 NM_172229 Homo sapiens kringle containing transmembrane protein 2 (KREMEN2), tran
 NM_172230 Homo sapiens synovial apoptosis inhibitor 1, synoviolin (SYVN1), transcript v
 NM_172231 Homo sapiens splicing factor 4 (SF4), transcript variant a, mRNA
 NM_172232 Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 5 (ABC/
 NM_172234 Homo sapiens interleukin 17 receptor B (IL17RB), transcript variant 2, mRNA
 NM_172236 Homo sapiens protein C-fucosyltransferase 1 (POFUT1), transcript variant 2,
 NM_172238 Homo sapiens transcription factor AP-2 beta (activating enhancer binding pr
 NM_172239 Homo sapiens exonuclease GOR (GOR), mRNA
 NM_172240 Homo sapiens TUWID12 (TUWID12), mRNA
 NM_172241 Homo sapiens cutaneous T-cell lymphoma-associated antigen 1 (CTAGE1),
 NM_172242 Homo sapiens sperm associated antigen 6 (SPAG6), transcript variant 2, mF
 NM_172244 Homo sapiens sarcoglycan, delta (35kDa dystrophin-associated glycoprotein
 NM_172245 Homo sapiens colony stimulating factor 2 receptor, alpha, low-affinity (granul
 NM_172246 Homo sapiens colony stimulating factor 2 receptor, alpha, low-affinity (granul
 NM_172247 Homo sapiens colony stimulating factor 2 receptor, alpha, low-affinity (granul
 NM_172248 Homo sapiens colony stimulating factor 2 receptor, alpha, low-affinity (granul
 NM_172249 Homo sapiens colony stimulating factor 2 receptor, alpha, low-affinity (granul
 NM_172250 Homo sapiens methylmalonic aciduria (cobalamin deficiency) type A (MMAA/
 NM_172251 Homo sapiens mitochondrial ribosomal protein L54 (MRPL54), nuclear gene
 NM_172311 Homo sapiens stoned B/TFIIA-alpha/beta-like factor (SALF), mRNA
 NM_172312 Homo sapiens sperm associated antigen 8 (SPAG8), transcript variant 2, mF
 NM_172313 Homo sapiens colony stimulating factor 3 receptor (granulocyte) (CSF3R), tr
 NM_172314 Homo sapiens interleukin 17E (IL17E), transcript variant 2, mRNA
 NM_172315 Homo sapiens Mels1, myeloid ecotropic viral integration site 1 homolog 2 (m
 NM_172316 Homo sapiens Mels1, myeloid ecotropic viral integration site 1 homolog 2 (m
 NM_172318 Homo sapiens potassium voltage-gated channel, subfamily G, member 1 (KC
 NM_172337 Homo sapiens orthodenticle homolog 2 (Drosophila) (OTX2), transcript varia
 NM_172341 Homo sapiens presenilin enhancer 2 (PEN2), mRNA
 NM_172343 Homo sapiens interleukin 17F (IL17F), transcript variant 2, mRNA
 NM_172344 Homo sapiens potassium voltage-gated channel, subfamily G, member 3 (KC
 NM_172345 Homo sapiens sperm associated antigen 9 (SPAG9), transcript variant 2, mF
 NM_172346 Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 6 (ABC/
 NM_172347 Homo sapiens potassium voltage-gated channel, subfamily G, member 4 (KC
 NM_172348 Homo sapiens interleukin 4 (IL4), transcript variant 2, mRNA
 NM_172349 Homo sapiens nuclear receptor binding SET domain protein 1 (NSD1), trans
 NM_172350 Homo sapiens membrane cofactor protein (CD46, trophoblast-lymphocyte cr
 NM_172351 Homo sapiens membrane cofactor protein (CD46, trophoblast-lymphocyte cr
 NM_172352 Homo sapiens membrane cofactor protein (CD46, trophoblast-lymphocyte cr
 NM_172353 Homo sapiens membrane cofactor protein (CD46, trophoblast-lymphocyte cr
 NM_172354 Homo sapiens membrane cofactor protein (CD46, trophoblast-lymphocyte cr
 NM_172355 Homo sapiens membrane cofactor protein (CD46, trophoblast-lymphocyte cr
 NM_172356 Homo sapiens membrane cofactor protein (CD46, trophoblast-lymphocyte cr

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NM_172357 Homo sapiens membrane cofactor protein (CD46, trophoblast-lymphocyte cr
 NM_172358 Homo sapiens membrane cofactor protein (CD46, trophoblast-lymphocyte cr
 NM_172359 Homo sapiens membrane cofactor protein (CD46, trophoblast-lymphocyte cr
 NM_172360 Homo sapiens membrane cofactor protein (CD46, trophoblast-lymphocyte cr
 NM_172361 Homo sapiens membrane cofactor protein (CD46, trophoblast-lymphocyte cr
 NM_172362 Homo sapiens potassium voltage-gated channel, subfamily H (eag-related), i
 NM_172364 Homo sapiens calcium channel, voltage-dependent, alpha 2/delta subunit 4 (I
 NM_172365 Homo sapiens chromosome 14 open reading frame 50 (C14orf50), mRNA
 NM_172366 Homo sapiens F-box protein 16 (FBXO16), mRNA
 NM_172367 Homo sapiens tumor suppressor candidate 5 (TUSC5), mRNA
 NM_172369 Homo sapiens complement component 1, q subcomponent, gamma polypep
 NM_172370 Homo sapiens D-amino acid oxidase activator (DAOA), mRNA
 NM_172373 Homo sapiens E74-like factor 1 (ets domain transcription factor) (ELF1), mRl
 NM_172374 Homo sapiens interleukin 4 induced 1 (IL4I1), transcript variant 2, mRNA
 NM_172375 Homo sapiens potassium voltage-gated channel, subfamily H (eag-related), i
 NM_172376 Homo sapiens potassium voltage-gated channel, subfamily H (eag-related), i
 NM_172377 Homo sapiens cancer/testis antigen 2 (CTAG2), transcript variant 1, mRNA
 NM_172386 Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 9 (ABC
 NM_172387 Homo sapiens nuclear factor of activated T-cells, cytoplasmic, calcineurin-de
 NM_172388 Homo sapiens nuclear factor of activated T-cells, cytoplasmic, calcineurin-de
 NM_172389 Homo sapiens nuclear factor of activated T-cells, cytoplasmic, calcineurin-de
 NM_172390 Homo sapiens nuclear factor of activated T-cells, cytoplasmic, calcineurin-de
 NM_173039 Homo sapiens aquaporin 11 (AQP11), mRNA
 NM_173042 Homo sapiens Interleukin 18 binding protein (IL18BP), transcript variant A, m
 NM_173043 Homo sapiens Interleukin 18 binding protein (IL18BP), transcript variant B, m
 NM_173044 Homo sapiens Interleukin 18 binding protein (IL18BP), transcript variant D, m
 NM_173050 Homo sapiens signal peptide, CUB domain, EGF-like 1 (SCUBE1), mRNA
 NM_173054 Homo sapiens reelin (RELN), transcript variant 2, mRNA
 NM_173055 Homo sapiens zonadhesin (ZAN), transcript variant 1, mRNA
 NM_173056 Homo sapiens zonadhesin (ZAN), transcript variant 2, mRNA
 NM_173057 Homo sapiens zonadhesin (ZAN), transcript variant 4, mRNA
 NM_173058 Homo sapiens zonadhesin (ZAN), transcript variant 5, mRNA
 NM_173059 Homo sapiens zonadhesin (ZAN), transcript variant 6, mRNA
 NM_173060 Homo sapiens calpastatin (CAST), transcript variant 2, mRNA
 NM_173061 Homo sapiens calpastatin (CAST), transcript variant 3, mRNA
 NM_173062 Homo sapiens calpastatin (CAST), transcript variant 4, mRNA
 NM_173064 Homo sapiens Interleukin 28 receptor, alpha (Interferon, lambda receptor) (IL
 NM_173065 Homo sapiens Interleukin 28 receptor, alpha (Interferon, lambda receptor) (IL
 NM_173073 Homo sapiens solute carrier family 35, member C2 (SLC35C2), transcript va
 NM_173074 Homo sapiens phosphatidylinositol glycan, class F (PIGF), transcript variant
 NM_173075 Homo sapiens amyloid beta (A4) precursor protein-binding, family B, membe
 NM_173076 Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 12 (ABC
 NM_173077 Homo sapiens carboxypeptidase O (CPO), mRNA
 NM_173078 Homo sapiens SLIT and NTRK-like family, member 4 (SLITRK4), mRNA
 NM_173079 Homo sapiens RUN domain containing 1 (RUNDC1), mRNA
 NM_173080 Homo sapiens small proline rich protein 4 (SPRR4), mRNA
 NM_173081 Homo sapiens armadillo repeat containing 3 (ARMC3), mRNA
 NM_173082 Homo sapiens SNF2 histone linker PHD RING helicase (SHPRH), mRNA
 NM_173083 Homo sapiens TUDOR gene similar (TGS), mRNA
 NM_173084 Homo sapiens tripartite motif-containing 59 (TRIM59), mRNA
 NM_173086 Homo sapiens keratin 6E (KRT6E), mRNA
 NM_173087 Homo sapiens calpain 3, (p94) (CAPN3), transcript variant 3, mRNA
 NM_173088 Homo sapiens calpain 3, (p94) (CAPN3), transcript variant 4, mRNA
 NM_173089 Homo sapiens calpain 3, (p94) (CAPN3), transcript variant 5, mRNA
 NM_173090 Homo sapiens calpain 3, (p94) (CAPN3), transcript variant 6, mRNA
 NM_173091 Homo sapiens nuclear factor of activated T-cells, cytoplasmic, calcineurin-de
 NM_173092 Homo sapiens potassium voltage-gated channel, subfamily H (eag-related), i

NM_173156 Homo sapiens chromosome 1 open reading frame 16 (C1orf16), transcript va
 NM_173157 Homo sapiens nuclear receptor subfamily 4, group A, member 1 (NR4A1), tr
 NM_173158 Homo sapiens nuclear receptor subfamily 4, group A, member 1 (NR4A1), tr
 NM_173159 Homo sapiens neuronal PAS domain protein 3 (NPAS3), mRNA
 NM_173160 Homo sapiens FXRD domain containing ion transport regulator 4 (FXRD4), tr
 NM_173161 Homo sapiens interleukin 1 family, member 10 (theta) (IL1F10), transcript va
 NM_173162 Homo sapiens potassium voltage-gated channel, subfamily H (eag-related), I
 NM_173163 Homo sapiens nuclear factor of activated T-cells, cytoplasmic, calcineurin-de
 NM_173164 Homo sapiens nuclear factor of activated T-cells, cytoplasmic, calcineurin-de
 NM_173165 Homo sapiens nuclear factor of activated T-cells, cytoplasmic, calcineurin-de
 NM_173167 Homo sapiens cardiomyopathy associated 4 (CMYA4), mRNA
 NM_173170 Homo sapiens interleukin 1 family, member 5 (delta) (IL1F5), transcript varia
 NM_173171 Homo sapiens nuclear receptor subfamily 4, group A, member 2 (NR4A2), tr
 NM_173172 Homo sapiens nuclear receptor subfamily 4, group A, member 2 (NR4A2), tr
 NM_173173 Homo sapiens nuclear receptor subfamily 4, group A, member 2 (NR4A2), tr
 NM_173174 Homo sapiens PTK2B protein tyrosine kinase 2 beta (PTK2B), transcript vari
 NM_173175 Homo sapiens PTK2B protein tyrosine kinase 2 beta (PTK2B), transcript vari
 NM_173176 Homo sapiens PTK2B protein tyrosine kinase 2 beta (PTK2B), transcript vari
 NM_173177 Homo sapiens nuclear DNA-binding protein (C1D), transcript variant 2, mRN
 NM_173178 Homo sapiens interleukin 1 family, member 8 (eta) (IL1F8), transcript variant
 NM_173179 Homo sapiens solute carrier family 35, member C2 (SLC35C2), transcript va
 NM_173191 Homo sapiens Kv channel interacting protein 2 (KCINIP2), transcript variant 2
 NM_173192 Homo sapiens Kv channel interacting protein 2 (KCINIP2), transcript variant 3
 NM_173193 Homo sapiens Kv channel interacting protein 2 (KCINIP2), transcript variant 4
 NM_173194 Homo sapiens Kv channel interacting protein 2 (KCINIP2), transcript variant 5
 NM_173195 Homo sapiens Kv channel interacting protein 2 (KCINIP2), transcript variant 6
 NM_173197 Homo sapiens Kv channel interacting protein 2 (KCINIP2), transcript variant 7
 NM_173198 Homo sapiens nuclear receptor subfamily 4, group A, member 3 (NR4A3), tr
 NM_173199 Homo sapiens nuclear receptor subfamily 4, group A, member 3 (NR4A3), tr
 NM_173200 Homo sapiens nuclear receptor subfamily 4, group A, member 3 (NR4A3), tr
 NM_173201 Homo sapiens ATPase, Ca⁺⁺ transporting, cardiac muscle, fast twitch 1 (ATP
 NM_173202 Homo sapiens interleukin 1 family, member 7 (zeta) (IL1F7), transcript varia
 NM_173203 Homo sapiens interleukin 1 family, member 7 (zeta) (IL1F7), transcript varia
 NM_173204 Homo sapiens interleukin 1 family, member 7 (zeta) (IL1F7), transcript varia
 NM_173205 Homo sapiens interleukin 1 family, member 7 (zeta) (IL1F7), transcript varia
 NM_173206 Homo sapiens protein inhibitor of activated STAT, 2 (PIAS2), transcript varia
 NM_173207 Homo sapiens TGFB-induced factor (TALE family homeobox) (TGIF), transcr
 NM_173208 Homo sapiens TGFB-induced factor (TALE family homeobox) (TGIF), transcr
 NM_173209 Homo sapiens TGFB-induced factor (TALE family homeobox) (TGIF), transcr
 NM_173210 Homo sapiens TGFB-induced factor (TALE family homeobox) (TGIF), transcr
 NM_173211 Homo sapiens TGFB-induced factor (TALE family homeobox) (TGIF), transcr
 NM_173213 Homo sapiens keratin 23 (histone deacetylase inducible) (KRT23), transcript
 NM_173214 Homo sapiens nuclear factor of activated T-cells 5, tonicity-responsive (NFA
 NM_173215 Homo sapiens nuclear factor of activated T-cells 5, tonicity-responsive (NFA
 NM_173216 Homo sapiens sialyltransferase 1 (beta-galactoside alpha-2,6-sialyltransferase)
 NM_173217 Homo sapiens sialyltransferase 1 (beta-galactoside alpha-2,6-sialyltransferase)
 NM_173341 Homo sapiens PHD finger protein 7 (PHF7), transcript variant 2, mRNA
 NM_173342 Homo sapiens Kv channel interacting protein 2 (KCINIP2), transcript variant 8
 NM_173343 Homo sapiens interleukin 1 receptor, type II (IL1R2), transcript variant 2, mR
 NM_173344 Homo sapiens sialyltransferase 4A (beta-galactoside alpha-2,3-sialyltransferase)
 NM_173351 Homo sapiens olfactory receptor, family 6, subfamily B, member 3 (OR6B3),
 NM_173352 Homo sapiens keratin 5b (K5B), mRNA
 NM_173353 Homo sapiens tryptophan hydroxylase 2 (TPH2), mRNA
 NM_173354 Homo sapiens SNF1-like kinase (SNF1LK), mRNA
 NM_173355 Homo sapiens uridine phosphorylase 2 (UPP2), mRNA
 NM_173357 Homo sapiens synovial sarcoma, X breakpoint 6 (SSX6), mRNA
 NM_173358 Homo sapiens synovial sarcoma, X breakpoint 7 (SSX7), mRNA

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NM_173360 Homo sapiens spermatogenesis associated 9 (SPATA9), transcript variant 2.
 NM_173362 Homo sapiens LOC317671 (LOC317671), mRNA
 NM_173452 Homo sapiens ficolin (collagen/fibrinogen domain containing) 3 (Hakata antigen)
 NM_173454 Homo sapiens phosphodiesterase 8A (PDE8A), transcript variant 2, mRNA
 NM_173455 Homo sapiens phosphodiesterase 8A (PDE8A), transcript variant 3, mRNA
 NM_173456 Homo sapiens phosphodiesterase 8A (PDE8A), transcript variant 4, mRNA
 NM_173457 Homo sapiens phosphodiesterase 8A (PDE8A), transcript variant 5, mRNA
 NM_173459 Homo sapiens interleukin 1 receptor-like 1 (IL1RL1), transcript variant 3, mRNA
 NM_173460 Homo sapiens defensin, beta 119 (DEFB119), transcript variant 2, mRNA
 NM_173462 Homo sapiens papilin, proteoglycan-like sulfated glycoprotein (PAPLN), mRNA
 NM_173463 Homo sapiens hypothetical protein DKFZp761B107 (DKFZp761B107), mRNA
 NM_173464 Homo sapiens I(3)mbt-like 4 (Drosophila) (L3MBTL4), mRNA
 NM_173465 Homo sapiens collagen, type XXIII, alpha 1 (COL23A1), mRNA
 NM_173466 Homo sapiens hypothetical protein DKFZp434P055 (DKFZp434P055), mRNA
 NM_173467 Homo sapiens malonyl-CoA:acyl carrier protein transacylase (malonyltransferase)
 NM_173468 Homo sapiens MOB1, Mps One Binder kinase activator-like 1A (yeast) (MOB1)
 NM_173469 Homo sapiens hypothetical protein LOC92912 (LOC92912), mRNA
 NM_173470 Homo sapiens transmembrane protein 32 (TMEM32), mRNA
 NM_173471 Homo sapiens solute carrier family 25, member 28 (SLC25A26), mRNA
 NM_173472 Homo sapiens hypothetical protein MGC40179 (MGC40179), mRNA
 NM_173473 Homo sapiens chromosome 10 open reading frame 104 (C10orf104), mRNA
 NM_173474 Homo sapiens N-terminal asparagine amidase (NTAN1), mRNA
 NM_173475 Homo sapiens hypothetical protein MGC48972 (MGC48972), mRNA
 NM_173476 Homo sapiens hypothetical protein FLJ34512 (FLJ34512), mRNA
 NM_173477 Homo sapiens Usher syndrome 1G (autosomal recessive) (USH1G), mRNA
 NM_173478 Homo sapiens hypothetical protein FLJ40137 (FLJ40137), mRNA
 NM_173479 Homo sapiens hypothetical protein LOC126248 (LOC126248), mRNA
 NM_173480 Homo sapiens hypothetical protein LOC126295 (LOC126295), mRNA
 NM_173481 Homo sapiens chromosome 19 open reading frame 21 (C19orf21), mRNA
 NM_173482 Homo sapiens hypothetical protein FLJ40365 (FLJ40365), mRNA
 NM_173483 Homo sapiens hypothetical protein FLJ39501 (FLJ39501), mRNA
 NM_173484 Homo sapiens hypothetical protein FLJ40160 (FLJ40160), mRNA
 NM_173485 Homo sapiens chromosome 20 open reading frame 17 (C20orf17), mRNA
 NM_173486 Homo sapiens hypothetical protein FLJ40298 (FLJ40298), mRNA
 NM_173487 Homo sapiens hypothetical protein LOC132321 (LOC132321), mRNA
 NM_173488 Homo sapiens solute carrier organic anion transporter family, member 6A1 (SLC6A1)
 NM_173489 Homo sapiens hypothetical protein FLJ40243 (FLJ40243), mRNA
 NM_173490 Homo sapiens hypothetical protein LOC134285 (LOC134285), mRNA
 NM_173491 Homo sapiens LSM11, U7 small nuclear RNA associated (LSM11), mRNA
 NM_173492 Homo sapiens phosphatidylinositol-4-phosphate 5-kinase-like 1 (PIP5KL1), mRNA
 NM_173493 Homo sapiens PAS domain containing protein 1 (PASD1), mRNA
 NM_173494 Homo sapiens hypothetical protein MGC35261 (MGC35261), mRNA
 NM_173495 Homo sapiens hypothetical protein FLJ30296 (FLJ30296), mRNA
 NM_173496 Homo sapiens membrane protein, palmitoylated 7 (MAGUK p55 subfamily member 7)
 NM_173497 Homo sapiens HECT domain containing 2 (HECTD2), transcript variant 2, mRNA
 NM_173499 Homo sapiens spermatogenesis-related protein 8 (MGC4294), mRNA
 NM_173500 Homo sapiens tau tubulin kinase 2 (TTBK2), mRNA
 NM_173501 Homo sapiens hypothetical protein LOC146174 (LOC146174), mRNA
 NM_173502 Homo sapiens hypothetical protein FLJ90661 (FLJ90661), mRNA
 NM_173503 Homo sapiens hypothetical protein FLJ25818 (FLJ25818), mRNA
 NM_173505 Homo sapiens ankyrin repeat domain 29 (ANKRD29), mRNA
 NM_173506 Homo sapiens hypothetical protein MGC42718 (MGC42718), mRNA
 NM_173507 Homo sapiens hypothetical protein FLJ37118 (FLJ37118), mRNA
 NM_173508 Homo sapiens solute carrier family 35, member F3 (SLC35F3), mRNA
 NM_173509 Homo sapiens hypothetical protein MGC16664 (MGC16664), mRNA
 NM_173510 Homo sapiens hypothetical protein FLJ33814 (FLJ33814), mRNA
 NM_173511 Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region,

NM_173512 Homo sapiens hypothetical protein FLJ39822 (FLJ39822), mRNA
 NM_173513 Homo sapiens hypothetical protein MGC43122 (MGC43122), mRNA
 NM_173514 Homo sapiens hypothetical protein FLJ90709 (FLJ90709), mRNA
 NM_173515 Homo sapiens membrane associated guanylate kinase interacting protein-lik
 NM_173516 Homo sapiens poly(A)-specific ribonuclease (PARN)-like domain containing '
 NM_173517 Homo sapiens vitamin K epoxide reductase complex, subunit 1-like 1 (VKOR
 NM_173518 Homo sapiens hypothetical protein FLJ25692 (FLJ25692), mRNA
 NM_173519 Homo sapiens hypothetical protein MGC34646 (MGC34646), mRNA
 NM_173521 Homo sapiens chromosome 9 open reading frame 84 (C9orf84), mRNA
 NM_173522 Homo sapiens hypothetical protein FLJ36576 (FLJ36576), mRNA
 NM_173523 Homo sapiens melanoma antigen, family B, 6 (MAGEB6), mRNA
 NM_173524 Homo sapiens chromosome 10 open reading frame 64 (C10orf64), mRNA
 NM_173525 Homo sapiens hypothetical protein MGC34805 (MGC34805), mRNA
 NM_173526 Homo sapiens chromosome 14 open reading frame 54 (C14orf54), mRNA
 NM_173527 Homo sapiens hypothetical protein FLJ38964 (FLJ38964), mRNA
 NM_173528 Homo sapiens chromosome 15 open reading frame 26 (C15orf26), mRNA
 NM_173529 Homo sapiens hypothetical protein MGC33382 (MGC33382), mRNA
 NM_173530 Homo sapiens zinc finger protein 610 (ZNF610), mRNA
 NM_173531 Homo sapiens zinc finger protein 100 (ZNF100), mRNA
 NM_173532 Homo sapiens hypothetical protein FLJ35838 (FLJ35838), mRNA
 NM_173533 Homo sapiens tudor domain containing 5 (TDRD5), mRNA
 NM_173535 Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain
 NM_173536 Homo sapiens gamma-aminobutyric acid (GABA) A receptor, gamma 1 (GAE
 NM_173537 Homo sapiens GTF2l repeat domain containing 2 (GTF2IRD2), mRNA
 NM_173538 Homo sapiens hypothetical protein FLJ35802 (FLJ35802), mRNA
 NM_173539 Homo sapiens zinc finger protein 596 (ZNF596), mRNA
 NM_173540 Homo sapiens fucosyltransferase 11 (alpha (1,3) fucosyltransferase) (FUT11
 NM_173541 Homo sapiens chromosome 10 open reading frame 91 (C10orf91), mRNA
 NM_173542 Homo sapiens hypothetical protein LOC196463 (LOC196463), mRNA
 NM_173543 Homo sapiens hypothetical protein FLJ32844 (FLJ32844), mRNA
 NM_173544 Homo sapiens B-cell novel protein 1 (BCNP1), mRNA
 NM_173545 Homo sapiens chromosome 2 open reading frame 13 (C2orf13), mRNA
 NM_173546 Homo sapiens hypothetical protein MGC35097 (MGC35097), mRNA
 NM_173547 Homo sapiens hypothetical protein LOC201292 (LOC201292), mRNA
 NM_173548 Homo sapiens zinc finger protein 584 (ZNF584), mRNA
 NM_173549 Homo sapiens hypothetical protein FLJ39553 (FLJ39553), mRNA
 NM_173550 Homo sapiens chromosome 9 open reading frame 93 (C9orf93), mRNA
 NM_173551 Homo sapiens sterile alpha motif domain containing 6 (SAMD6), mRNA
 NM_173552 Homo sapiens hypothetical protein MGC33365 (MGC33365), mRNA
 NM_173553 Homo sapiens hypothetical protein FLJ25801 (FLJ25801), mRNA
 NM_173554 Homo sapiens chromosome 10 open reading frame 107 (C10orf107), mRNA
 NM_173555 Homo sapiens trypsin domain containing 1 (TYSND1), mRNA
 NM_173556 Homo sapiens hypothetical protein MGC34732 (MGC34732), mRNA
 NM_173557 Homo sapiens ring finger protein 152 (RNF152), mRNA
 NM_173558 Homo sapiens FGD1 family, member 2 (FGD2), mRNA
 NM_173559 Homo sapiens hypothetical protein FLJ25791 (FLJ25791), mRNA
 NM_173560 Homo sapiens regulatory factor X domain containing 1 (RFXDC1), mRNA
 NM_173561 Homo sapiens unc-5 homolog C (C. elegans)-like (UNC5CL), mRNA
 NM_173562 Homo sapiens chromosome 6 open reading frame 69 (C6orf69), mRNA
 NM_173563 Homo sapiens chromosome 6 open reading frame 146 (C6orf146), mRNA
 NM_173564 Homo sapiens hypothetical protein FLJ37538 (FLJ37538), mRNA
 NM_173565 Homo sapiens hypothetical protein LOC222967 (LOC222967), mRNA
 NM_173566 Homo sapiens hypothetical protein MGC50372 (MGC50372), mRNA
 NM_173567 Homo sapiens abhydrolase domain containing 7 (ABHD7), mRNA
 NM_173568 Homo sapiens uromodulin-like 1 (UMODL1), mRNA
 NM_173570 Homo sapiens zinc finger, DHHC domain containing 23 (ZDHHC23), mRNA
 NM_173571 Homo sapiens hypothetical protein LOC255313 (LOC255313), mRNA

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NM_173572 Homo sapiens chromosome 10 open reading frame 93 (C10orf93), mRNA
NM_173573 Homo sapiens hypothetical protein MGC35138 (MGC35138), mRNA
NM_173574 Homo sapiens hypothetical protein MGC33414 (MGC33414), mRNA
NM_173575 Homo sapiens serine/threonine kinase 32C (STK32C), mRNA
NM_173576 Homo sapiens chromosome 10 open reading frame 48 (C10orf48), mRNA
NM_173578 Homo sapiens hypothetical protein FLJ90834 (FLJ90834), mRNA
NM_173579 Homo sapiens hypothetical protein FLJ40224 (FLJ40224), mRNA
NM_173580 Homo sapiens hypothetical protein FLJ39058 (FLJ39058), mRNA
NM_173581 Homo sapiens hypothetical protein FLJ90231 (FLJ90231), mRNA
NM_173582 Homo sapiens phosphoglucomutase 2-like 1 (PGM2L1), mRNA
NM_173583 Homo sapiens hypothetical protein FLJ33790 (FLJ33790), mRNA
NM_173584 Homo sapiens hypothetical protein MGC45840 (MGC45840), mRNA
NM_173586 Homo sapiens hypothetical protein MGC34821 (MGC34821), mRNA
NM_173587 Homo sapiens REST corepressor 2 (RCOR2), mRNA
NM_173588 Homo sapiens hypothetical protein FLJ37794 (FLJ37794), mRNA
NM_173589 Homo sapiens hypothetical protein FLJ35709 (FLJ35709), mRNA
NM_173590 Homo sapiens hypothetical protein FLJ36102 (FLJ36102), mRNA
NM_173591 Homo sapiens hypothetical protein FLJ90579 (FLJ90579), mRNA
NM_173593 Homo sapiens beta 1,4-N-acetylgalactosaminyltransferase-transferase-III (B-
NM_173596 Homo sapiens solute carrier family 39 (metal ion transporter), member 5 (SL-
NM_173597 Homo sapiens hypothetical protein FLJ37567 (FLJ37567), mRNA
NM_173598 Homo sapiens kinase suppressor of Ras-2 (KSR2), mRNA
NM_173599 Homo sapiens hypothetical protein FLJ40126 (FLJ40126), mRNA
NM_173605 Homo sapiens potassium channel regulator (KCNRG), mRNA
NM_173607 Homo sapiens chromosome 14 open reading frame 24 (C14orf24), mRNA
NM_173608 Homo sapiens chromosome 14 open reading frame 80 (C14orf80), mRNA
NM_173609 Homo sapiens chromosome 15 open reading frame 21 (C15orf21), mRNA
NM_173610 Homo sapiens hypothetical protein FLJ33788 (FLJ33788), mRNA
NM_173611 Homo sapiens hypothetical protein FLJ38426 (FLJ38426), mRNA
NM_173613 Homo sapiens hypothetical protein FLJ35785 (FLJ35785), mRNA
NM_173614 Homo sapiens hypothetical protein LOC283820 (LOC283820), mRNA
NM_173616 Homo sapiens hypothetical protein FLJ35894 (FLJ35894), mRNA
NM_173617 Homo sapiens hypothetical protein FLJ36701 (FLJ36701), mRNA
NM_173618 Homo sapiens hypothetical protein FLJ90652 (FLJ90652), mRNA
NM_173619 Homo sapiens hypothetical protein MGC34781 (MGC34781), mRNA
NM_173620 Homo sapiens hypothetical protein FLJ23825 (FLJ23825), mRNA
NM_173621 Homo sapiens hypothetical protein FLJ34790 (FLJ34790), mRNA
NM_173622 Homo sapiens hypothetical protein FLJ36674 (FLJ36674), mRNA
NM_173623 Homo sapiens hypothetical protein FLJ35808 (FLJ35808), mRNA
NM_173624 Homo sapiens hypothetical protein FLJ40504 (FLJ40504), mRNA
NM_173625 Homo sapiens hypothetical protein FLJ39647 (FLJ39647), mRNA
NM_173626 Homo sapiens solute carrier family 26, member 11 (SLC28A11), mRNA
NM_173627 Homo sapiens hypothetical protein FLJ35220 (FLJ35220), mRNA
NM_173628 Homo sapiens hypothetical protein FLJ40457 (FLJ40457), mRNA
NM_173629 Homo sapiens chromosome 18 open reading frame 26 (C18orf26), mRNA
NM_173630 Homo sapiens rotatin (RTTN), mRNA
NM_173631 Homo sapiens zinc finger protein 547 (ZNF547), mRNA
NM_173632 Homo sapiens hypothetical protein FLJ38288 (FLJ38288), mRNA
NM_173633 Homo sapiens hypothetical protein FLJ90805 (FLJ90805), mRNA
NM_173635 Homo sapiens hypothetical protein FLJ40235 (FLJ40235), mRNA
NM_173636 Homo sapiens chromosome 19 open reading frame 14 (C19orf14), mRNA
NM_173637 Homo sapiens hypothetical protein MGC34725 (MGC34725), mRNA
NM_173638 Homo sapiens hypothetical protein MGC8902 (MGC8902), mRNA
NM_173639 Homo sapiens hypothetical protein FLJ35976 (FLJ35976), mRNA
NM_173640 Homo sapiens likely ortholog of mouse roof plate-specific spondin (R-spondi
NM_173641 Homo sapiens hypothetical protein FLJ33655 (FLJ33655), mRNA
NM_173642 Homo sapiens hypothetical protein MGC47816 (MGC47816), mRNA

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NM_173643 Homo sapiens hypothetical protein DKFZp547G0215 (DKFZp547G0215), mi
 NM_173644 Homo sapiens hypothetical protein FLJ33860 (FLJ33860), mRNA
 NM_173645 Homo sapiens hypothetical protein FLJ37357 (FLJ37357), mRNA
 NM_173646 Homo sapiens hypothetical protein FLJ39660 (FLJ39660), mRNA
 NM_173647 Homo sapiens ring finger protein 149 (RNF149), mRNA
 NM_173648 Homo sapiens hypothetical protein FLJ39502 (FLJ39502), mRNA
 NM_173649 Homo sapiens hypothetical protein FLJ40172 (FLJ40172), mRNA
 NM_173650 Homo sapiens DnaJ (Hsp40) homolog, subfamily C, member 5 gamma (DN/
 NM_173651 Homo sapiens fibrous sheath interacting protein 2 (FSIP2), mRNA
 NM_173652 Homo sapiens hypothetical protein MGC34824 (MGC34824), mRNA
 NM_173653 Homo sapiens solute carrier family 9 (sodium/hydrogen exchanger), isoform
 NM_173654 Homo sapiens hypothetical protein MGC34132 (MGC34132), mRNA
 NM_173656 Homo sapiens zinc finger protein 619 (ZNF619), mRNA
 NM_173657 Homo sapiens hypothetical protein FLJ31139 (FLJ31139), mRNA
 NM_173658 Homo sapiens hypothetical protein FLJ36870 (FLJ36870), mRNA
 NM_173659 Homo sapiens hypothetical protein MGC29784 (MGC29784), mRNA
 NM_173660 Homo sapiens hypothetical protein FLJ33718 (FLJ33718), mRNA
 NM_173661 Homo sapiens hypothetical protein FLJ35424 (FLJ35424), mRNA
 NM_173662 Homo sapiens hypothetical protein LOC285533 (LOC285533), mRNA
 NM_173663 Homo sapiens NY-REN-7 antigen (NY-REN-7), mRNA
 NM_173664 Homo sapiens ADP-ribosylation factor-like 10A (ARL10A), mRNA
 NM_173665 Homo sapiens hypothetical protein MGC34713 (MGC34713), mRNA
 NM_173666 Homo sapiens hypothetical protein FLJ33977 (FLJ33977), mRNA
 NM_173667 Homo sapiens hypothetical protein FLJ37543 (FLJ37543), mRNA
 NM_173669 Homo sapiens hypothetical protein FLJ34047 (FLJ34047), mRNA
 NM_173670 Homo sapiens RGM domain family, member B (RGM), mRNA
 NM_173671 Homo sapiens hypothetical protein FLJ37396 (FLJ37396), mRNA
 NM_173672 Homo sapiens peptidylprolyl isomerase (cyclophilin)-like 6 (PPI6), mRNA
 NM_173673 Homo sapiens hypothetical protein FLJ34503 (FLJ34503), mRNA
 NM_173674 Homo sapiens discoidin, CUB and LCCL domain containing 1 (DCBLD1), mf
 NM_173675 Homo sapiens hypothetical protein FLJ33708 (FLJ33708), mRNA
 NM_173676 Homo sapiens patatin-like phospholipase domain containing 1 (PNPLA1), ml
 NM_173677 Homo sapiens hypothetical protein FLJ40852 (FLJ40852), mRNA
 NM_173678 Homo sapiens hypothetical protein FLJ40722 (FLJ40722), mRNA
 NM_173680 Homo sapiens hypothetical protein MGC33584 (MGC33584), mRNA
 NM_173682 Homo sapiens hypothetical protein FLJ40288 (FLJ40288), mRNA
 NM_173683 Homo sapiens chromosome 8 open reading frame 21 (C8orf21), mRNA
 NM_173685 Homo sapiens hypothetical protein FLJ32440 (FLJ32440), mRNA
 NM_173687 Homo sapiens hypothetical protein FLJ37131 (FLJ37131), mRNA
 NM_173688 Homo sapiens hypothetical protein FLJ39630 (FLJ39630), mRNA
 NM_173689 Homo sapiens crumbs homolog 2 (Drosophila) (CRB2), mRNA
 NM_173690 Homo sapiens chromosome 9 open reading frame 126 (C9orf126), mRNA
 NM_173691 Homo sapiens chromosome 9 open reading frame 75 (C9orf75), mRNA
 NM_173694 Homo sapiens ATPase, Class VI, type 11C (ATP11C), mRNA
 NM_173695 Homo sapiens hypothetical protein FLJ36601 (FLJ36601), mRNA
 NM_173698 Homo sapiens hypothetical protein FLJ37659 (FLJ37659), mRNA
 NM_173699 Homo sapiens hypothetical protein MGC33889 (MGC33889), mRNA
 NM_173700 Homo sapiens BCL6 co-repressor-like 2 (BCORL2), mRNA
 NM_173701 Homo sapiens tryptophanyl-IRNA synthetase (WARS), transcript variant 2, tr
 NM_173728 Homo sapiens Rho guanine nucleotide exchange factor (GEF) 15 (ARHGGEF/
 NM_173791 Homo sapiens PDZ domain containing 8 (PDZK8), mRNA
 NM_173793 Homo sapiens hypothetical protein LOC128977 (LOC128977), mRNA
 NM_173794 Homo sapiens FUN14 domain containing 1 (FUND1), mRNA
 NM_173795 Homo sapiens hypothetical protein FLJ32096 (FLJ32096), mRNA
 NM_173797 Homo sapiens PAP associated domain containing 4 (PAPD4), mRNA
 NM_173798 Homo sapiens zinc finger, CCHC domain containing 12 (ZCCHC12), mRNA
 NM_173799 Homo sapiens hypothetical protein FLJ39873 (FLJ39873), mRNA

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NM_173800 Homo sapiens laeverin (FLJ90650), mRNA
 NM_173801 Homo sapiens hypothetical protein FLJ36198 (FLJ36198), mRNA
 NM_173802 Homo sapiens hypothetical protein MGC50559 (MGC50559), mRNA
 NM_173803 Homo sapiens hypothetical protein FLJ39599 (FLJ39599), mRNA
 NM_173804 Homo sapiens hypothetical protein MGC30208 (MGC30208), mRNA
 NM_173805 Homo sapiens hypothetical protein FLJ38723 (FLJ38723), mRNA
 NM_173806 Homo sapiens hypothetical protein MGC50721 (MGC50721), mRNA
 NM_173807 Homo sapiens hypothetical protein MGC33370 (MGC33370), mRNA
 NM_173808 Homo sapiens neuronal growth regulator 1 (NEGR1), mRNA
 NM_173809 Homo sapiens biogenesis of lysosome-related organelles complex-1, subunit
 NM_173810 Homo sapiens hypothetical protein MGC29649 (MGC29649), mRNA
 NM_173811 Homo sapiens hypothetical protein FLJ32675 (FLJ32675), mRNA
 NM_173812 Homo sapiens hypothetical protein FLJ32949 (FLJ32949), mRNA
 NM_173813 Homo sapiens hypothetical protein FLJ34154 (FLJ34154), mRNA
 NM_173815 Homo sapiens hypothetical protein FLJ37464 (FLJ37464), mRNA
 NM_173821 Homo sapiens hypothetical protein FLJ33590 (FLJ33590), mRNA
 NM_173822 Homo sapiens hypothetical protein MGC39518 (MGC39518), mRNA
 NM_173824 Homo sapiens hypothetical protein MGC26717 (MGC26717), mRNA
 NM_173825 Homo sapiens RAB, member of RAS oncogene family-like 3 (RABL3), mRNA/
 NM_173826 Homo sapiens hypothetical protein DKFZp313N0621 (DKFZp313N0621), mf
 NM_173827 Homo sapiens hypothetical protein FLJ38991 (FLJ38991), mRNA
 NM_173828 Homo sapiens chromosome 5 open reading frame 16 (C5orf16), mRNA
 NM_173829 Homo sapiens hypothetical protein FLJ36754 (FLJ36754), mRNA
 NM_173830 Homo sapiens chromosome 6 open reading frame 182 (C6orf182), mRNA
 NM_173831 Homo sapiens hypothetical protein LOC286075 (LOC286075), mRNA
 NM_173832 Homo sapiens hypothetical protein FLJ38705 (FLJ38705), mRNA
 NM_173833 Homo sapiens hypothetical protein MGC45780 (MGC45780), mRNA
 NM_173834 Homo sapiens hypothetical protein MGC21416 (MGC21416), mRNA
 NM_173841 Homo sapiens interleukin 1 receptor antagonist (IL1RN), transcript variant 2,
 NM_173842 Homo sapiens interleukin 1 receptor antagonist (IL1RN), transcript variant 1,
 NM_173843 Homo sapiens interleukin 1 receptor antagonist (IL1RN), transcript variant 4,
 NM_173844 Homo sapiens mucosa associated lymphoid tissue lymphoma translocation c
 NM_173846 Homo sapiens chromosome 14 open reading frame 8 (C14orf8), mRNA
 NM_173847 Homo sapiens sperm acrosome associated 3 (SPACA3), mRNA
 NM_173848 Homo sapiens hypothetical protein LOC138046 (LOC138046), mRNA
 NM_173849 Homo sapiens goosecoid (GSC), mRNA
 NM_173850 Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1 anti
 NM_173851 Homo sapiens solute carrier family 30 (zinc transporter), member 8 (SLC30A
 NM_173852 Homo sapiens keratinocyte associated protein 2 (KRTCAP2), mRNA
 NM_173853 Homo sapiens keratinocyte associated protein 3 (KRTCAP3), mRNA
 NM_173854 Homo sapiens solute carrier family 41, member 1 (SLC41A1), mRNA
 NM_173855 Homo sapiens morn (LOC283385), mRNA
 NM_173856 Homo sapiens vomeronasal 1 receptor 2 (VN1R2), mRNA
 NM_173857 Homo sapiens vomeronasal 1 receptor 4 (VN1R4), mRNA
 NM_173858 Homo sapiens vomeronasal 1 receptor 5 (VN1R5), mRNA
 NM_173859 Homo sapiens breast cancer and salivary gland expression gene (BASE), ml
 NM_173860 Homo sapiens homeo box C12 (HOXC12), mRNA
 NM_173872 Homo sapiens chloride channel 3 (CLCN3), transcript variant e, mRNA
 NM_174855 Homo sapiens isocitrate dehydrogenase 3 (NAD+) beta (IDH3B), nuclear ge
 NM_174856 Homo sapiens isocitrate dehydrogenase 3 (NAD+) beta (IDH3B), nuclear ge
 NM_174858 Homo sapiens adenylate kinase 5 (AK5), transcript variant 1, mRNA
 NM_174869 Homo sapiens isocitrate dehydrogenase 3 (NAD+) gamma (IDH3G), nuclear
 NM_174871 Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase,
 NM_174872 Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 2 (P2RX2)
 NM_174873 Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 2 (P2RX2)
 NM_174878 Homo sapiens Usher syndrome 3A (USH3A), transcript variant 1, mRNA
 NM_174880 Homo sapiens Usher syndrome 3A (USH3A), transcript variant 3, mRNA

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NM_174881 Homo sapiens crumbs homolog 3 (Drosophila) (CRB3), transcript variant 3, r
 NM_174882 Homo sapiens crumbs homolog 3 (Drosophila) (CRB3), transcript variant 1, r
 NM_174886 Homo sapiens TGF β -induced factor (TALE family homeobox) (TGIF), transcr
 NM_174887 Homo sapiens intraflagellar transport protein IFT20 (LOC90410), mRNA
 NM_174889 Homo sapiens hypothetical protein LOC91942 (LOC91942), mRNA
 NM_174890 Homo sapiens AN1, ubiquitin-like, homolog (Xenopus laevis) (ANUBL1), mR
 NM_174891 Homo sapiens chromosome 14 open reading frame 79 (C14orf79), mRNA
 NM_174892 Homo sapiens triggering receptor expressed on myeloid cells 5 (TREM5), mR
 NM_174896 Homo sapiens hypothetical protein MGC24133 (MGC24133), mRNA
 NM_174897 Homo sapiens bactericidal/permeability-increasing protein-like 3 (BPII3), mF
 NM_174898 Homo sapiens hypothetical protein LOC129530 (LOC129530), mRNA
 NM_174899 Homo sapiens F-box protein 36 (FBXO36), mRNA
 NM_174900 Homo sapiens zinc finger protein 42 (ZFP42), mRNA
 NM_174901 Homo sapiens family with sequence similarity 9, member C (FAM9C), mRNA
 NM_174902 Homo sapiens hypothetical protein LOC143458 (LOC143458), mRNA
 NM_174905 Homo sapiens hypothetical protein LOC147965 (LOC147965), mRNA
 NM_174906 Homo sapiens hypothetical protein MGC39724 (MGC39724), mRNA
 NM_174907 Homo sapiens protein phosphatase 4, regulatory subunit 2 (PPP4R2), mRN/
 NM_174908 Homo sapiens chromosome 3 open reading frame 6 (C3orf6), transcript vari
 NM_174909 Homo sapiens hypothetical protein MGC23909 (MGC23909), mRNA
 NM_174910 Homo sapiens t-complex-associated-testis-expressed 3 (TCTE3), mRNA
 NM_174911 Homo sapiens breast cancer membrane protein 101 (NSE2), mRNA
 NM_174912 Homo sapiens hypothetical protein FLJ31204 (FLJ31204), mRNA
 NM_174913 Homo sapiens chromosome 14 open reading frame 21 (C14orf21), mRNA
 NM_174914 Homo sapiens hypothetical protein LOC167127 (LOC167127), mRNA
 NM_174916 Homo sapiens ubiquitin protein ligase E3 component n-recognin 1 (UBR1), n
 NM_174917 Homo sapiens hypothetical protein LOC197322 (LOC197322), mRNA
 NM_174918 Homo sapiens hypothetical protein LOC199675 (LOC199675), mRNA
 NM_174920 Homo sapiens hypothetical protein LOC201191 (LOC201191), mRNA
 NM_174921 Homo sapiens hypothetical protein LOC201895 (LOC201895), mRNA
 NM_174922 Homo sapiens aarF domain containing kinase 5 (ADCK5), mRNA
 NM_174923 Homo sapiens hypothetical protein MGC31967 (MGC31967), mRNA
 NM_174924 Homo sapiens hypothetical protein LOC204474 (LOC204474), mRNA
 NM_174925 Homo sapiens hypothetical protein LOC205251 (LOC205251), mRNA
 NM_174926 Homo sapiens hypothetical protein MGC17839 (MGC17839), mRNA
 NM_174927 Homo sapiens spergen-1 (SPAS1), mRNA
 NM_174928 Homo sapiens hypothetical protein LOC221143 (LOC221143), mRNA
 NM_174930 Homo sapiens postmeiotic segregation increased 2-like 5 (PMS2L5), mRNA
 NM_174931 Homo sapiens hypothetical protein FLJ38348 (FLJ38348), mRNA
 NM_174932 Homo sapiens bactericidal/permeability-increasing protein-like 2 (BPII2), mF
 NM_174933 Homo sapiens phytoey-CoA dioxygenase domain containing 1 (PHYHD1),
 NM_174934 Homo sapiens sodium channel, voltage-gated, type IV, beta (SCN4B), mRN/
 NM_174936 Homo sapiens proprotein convertase subtilisin/kexin type 9 (PCSK9), mRNA
 NM_174937 Homo sapiens transcription elongation regulator 1-like (TCERG1L), mRNA
 NM_174938 Homo sapiens FERM domain containing 3 (FRMD3), mRNA
 NM_174939 Homo sapiens hypothetical protein MGC39681 (MGC39681), mRNA
 NM_174940 Homo sapiens hypothetical protein LOC283232 (LOC283232), mRNA
 NM_174941 Homo sapiens scavenger receptor cysteine-rich type 1 protein M160 (M160),
 NM_174942 Homo sapiens growth arrest-specific 2 like 3 (GAS2L3), mRNA
 NM_174943 Homo sapiens hypothetical protein FLJ25976 (FLJ25976), mRNA
 NM_174944 Homo sapiens chromosome 14 open reading frame 20 (C14orf20), mRNA
 NM_174945 Homo sapiens zinc finger protein 575 (ZNF575), mRNA
 NM_174947 Homo sapiens chromosome 19 open reading frame 30 (C19orf30), mRNA
 NM_174950 Homo sapiens hypothetical protein FLJ30435 (FLJ30435), mRNA
 NM_174951 Homo sapiens family with sequence similarity 9, member A (FAM9A), mRNA
 NM_174952 Homo sapiens hypothetical protein MGC46496 (MGC46496), mRNA
 NM_174953 Homo sapiens ATPase, Ca⁺⁺ transporting, ubiquitous (ATP2A3), transcript v

NM_174954 Homo sapiens ATPase, Ca++ transporting, ubiquitous (ATP2A3), transcript v
 NM_174955 Homo sapiens ATPase, Ca++ transporting, ubiquitous (ATP2A3), transcript v
 NM_174956 Homo sapiens ATPase, Ca++ transporting, ubiquitous (ATP2A3), transcript v
 NM_174957 Homo sapiens ATPase, Ca++ transporting, ubiquitous (ATP2A3), transcript v
 NM_174958 Homo sapiens ATPase, Ca++ transporting, ubiquitous (ATP2A3), transcript v
 NM_174959 Homo sapiens hypothetical protein LOC136306 (LOC136306), mRNA
 NM_174961 Homo sapiens synovial sarcoma, X breakpoint 8 (SSX8), mRNA
 NM_174962 Homo sapiens synovial sarcoma, X breakpoint 9 (SSX9), mRNA
 NM_174963 Homo sapiens sialyltransferase 6 (N-acetylglucosaminide alpha 2,3-sialyltrans
 NM_174964 Homo sapiens sialyltransferase 6 (N-acetylglucosaminide alpha 2,3-sialyltrans
 NM_174965 Homo sapiens sialyltransferase 6 (N-acetylglucosaminide alpha 2,3-sialyltrans
 NM_174966 Homo sapiens sialyltransferase 6 (N-acetylglucosaminide alpha 2,3-sialyltrans
 NM_174967 Homo sapiens sialyltransferase 6 (N-acetylglucosaminide alpha 2,3-sialyltrans
 NM_174968 Homo sapiens sialyltransferase 6 (N-acetylglucosaminide alpha 2,3-sialyltrans
 NM_174969 Homo sapiens sialyltransferase 6 (N-acetylglucosaminide alpha 2,3-sialyltrans
 NM_174970 Homo sapiens sialyltransferase 6 (N-acetylglucosaminide alpha 2,3-sialyltrans
 NM_174971 Homo sapiens sialyltransferase 6 (N-acetylglucosaminide alpha 2,3-sialyltrans
 NM_174972 Homo sapiens sialyltransferase 6 (N-acetylglucosaminide alpha 2,3-sialyltrans
 NM_174975 Homo sapiens SEC14-like 3 (S. cerevisiae) (SEC14L3), mRNA
 NM_174976 Homo sapiens zinc finger, DHHC domain containing 22 (ZDHHC22), mRNA
 NM_174977 Homo sapiens SEC14-like 4 (S. cerevisiae) (SEC14L4), mRNA
 NM_174978 Homo sapiens chromosome 14 open reading frame 39 (C14orf39), mRNA
 NM_174980 Homo sapiens vomeronasal 1 receptor 3 (VN1R3), mRNA
 NM_174981 Homo sapiens ankyrin repeat domain 21 (ANKRD21), mRNA
 NM_174983 Homo sapiens chromosome 19 open reading frame 28 (C19orf28), mRNA
 NM_175038 Homo sapiens contactin 1 (CNTN1), transcript variant 2, mRNA
 NM_175039 Homo sapiens sialyltransferase 7D (alpha-N-acetylneuraminyl-2,3-beta-gala
 NM_175040 Homo sapiens sialyltransferase 7D (alpha-N-acetylneuraminyl-2,3-beta-gala
 NM_175047 Homo sapiens paired immunoglobulin-like type 2 receptor beta (PILRB), transc
 NM_175052 Homo sapiens sialyltransferase 8D (alpha-2, 8-polysialyltransferase) (SIAT8I
 NM_175053 Homo sapiens keratin 6 lrs4 (K6IRS4), mRNA
 NM_175054 Homo sapiens histone 4, H4 (HIST4H4), mRNA
 NM_175055 Homo sapiens histone 3, H2bb (HIST3H2BB), mRNA
 NM_175056 Homo sapiens hypothetical protein LOC131368 (LOC131368), mRNA
 NM_175057 Homo sapiens trace amine receptor 3 (TRAR3), mRNA
 NM_175058 Homo sapiens hypothetical protein LOC144100 (LOC144100), mRNA
 NM_175080 Homo sapiens chromosome 14 open reading frame 27 (C14orf27), mRNA
 NM_175081 Homo sapiens juxtaposed with another zinc finger gene 1 (JAZF1), mRNA
 NM_175082 Homo sapiens RasGEF domain family, member 1C (RASGEF1C), mRNA
 NM_175083 Homo sapiens hypothetical protein LOC284361 (LOC284361), transcript vari
 NM_175084 Homo sapiens Williams Beuren syndrome chromosome region 19 (WBSCR1
 NM_175085 Homo sapiens histone 2, H2ab (HIST2H2AB), mRNA
 NM_175086 Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 51 (DDX51), mRNA
 NM_175087 Homo sapiens trace amine receptor 4 (TRAR4), mRNA
 NM_175088 Homo sapiens keratin 6 lrs3 (K6IRS3), mRNA
 NM_175089 Homo sapiens aprataxin (APTX), transcript variant 2, mRNA
 NM_175071 Homo sapiens aprataxin (APTX), transcript variant 5, mRNA
 NM_175072 Homo sapiens aprataxin (APTX), transcript variant 3, mRNA
 NM_175073 Homo sapiens aprataxin (APTX), transcript variant 1, mRNA
 NM_175075 Homo sapiens hypothetical protein INMO1 (INMO1), mRNA
 NM_175077 Homo sapiens Src-like-adaptor 2 (SLA2), transcript variant 2, mRNA
 NM_175078 Homo sapiens keratin 1B (KRT1B), mRNA
 NM_175080 Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 5 (P2RX5)
 NM_175081 Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 5 (P2RX5)
 NM_175085 Homo sapiens phosphoribosylglycinamide formyltransferase, phosphoribosyl
 NM_175566 Homo sapiens contactin 5 (CNTN5), transcript variant 2, mRNA
 NM_175567 Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 4 (P2RX4)

NM_175568 Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 4 (P2RX4)
 NM_175569 Homo sapiens Xg blood group (pseudautosomal boundary-divided on the X
 NM_175571 Homo sapiens human immune associated nucleotide 6 (hIAN6), mRNA
 NM_175573 Homo sapiens adhesion regulating molecule 1 (ADRM1), transcript variant 2,
 NM_175575 Homo sapiens WFIKK-related protein (WFIKKRNP), mRNA
 NM_175605 Homo sapiens tetratricopeptide repeat domain 10 (TTC10), transcript variant
 NM_175607 Homo sapiens contactin 4 (CNTN4), transcript variant 1, mRNA
 NM_175609 Homo sapiens ADP-ribosylation factor GTPase activating protein 1 (ARFGAP
 NM_175610 Homo sapiens tight junction protein 1 (zona occludens 1) (TJP1), transcript v
 NM_175611 Homo sapiens glutamate receptor, ionotropic, kainate 1 (GRIK1), transcript v
 NM_175612 Homo sapiens contactin 4 (CNTN4), transcript variant 2, mRNA
 NM_175613 Homo sapiens contactin 4 (CNTN4), transcript variant 3, mRNA
 NM_175614 Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 11,
 NM_175616 Homo sapiens FIS (FIS), mRNA
 NM_175617 Homo sapiens metallothionein 1E (functional) (MT1E), mRNA
 NM_175619 Homo sapiens zygote arrest 1 (ZAR1), mRNA
 NM_175622 Homo sapiens metallothionein 1J (MT1J), mRNA
 NM_175623 Homo sapiens RAB3A interacting protein (rabn3) (RAB3IP), transcript variar
 NM_175624 Homo sapiens RAB3A interacting protein (rabn3) (RAB3IP), transcript variar
 NM_175625 Homo sapiens RAB3A interacting protein (rabn3) (RAB3IP), transcript variar
 NM_175626 Homo sapiens RAB3A interacting protein (rabn3) (RAB3IP), transcript variar
 NM_175627 Homo sapiens RAB3A interacting protein (rabn3) (RAB3IP), transcript variar
 NM_175629 Homo sapiens DNA (cytosine-5)-methyltransferase 3 alpha (DNMT3A), tran
 NM_175630 Homo sapiens DNA (cytosine-5)-methyltransferase 3 alpha (DNMT3A), tran
 NM_175634 Homo sapiens core-binding factor, runt domain, alpha subunit 2; translocat
 NM_175635 Homo sapiens core-binding factor, runt domain, alpha subunit 2; translocat
 NM_175636 Homo sapiens core-binding factor, runt domain, alpha subunit 2; translocat
 NM_175698 Homo sapiens synovial sarcoma, X breakpoint 2 (SSX2), transcript variant 2,
 NM_175709 Homo sapiens chromobox homolog 7 (CBX7), mRNA
 NM_175711 Homo sapiens synovial sarcoma, X breakpoint 3 (SSX3), transcript variant 2,
 NM_175719 Homo sapiens thyroid peroxidase (TPO), transcript variant 2, mRNA
 NM_175720 Homo sapiens thyroid peroxidase (TPO), transcript variant 3, mRNA
 NM_175721 Homo sapiens thyroid peroxidase (TPO), transcript variant 4, mRNA
 NM_175722 Homo sapiens thyroid peroxidase (TPO), transcript variant 5, mRNA
 NM_175723 Homo sapiens synovial sarcoma, X breakpoint 5 (SSX5), transcript variant 2,
 NM_175724 Homo sapiens interleukin 5 receptor, alpha (IL5RA), transcript variant 2, mRl
 NM_175725 Homo sapiens interleukin 5 receptor, alpha (IL5RA), transcript variant 3, mRl
 NM_175726 Homo sapiens interleukin 5 receptor, alpha (IL5RA), transcript variant 4, mRl
 NM_175727 Homo sapiens interleukin 5 receptor, alpha (IL5RA), transcript variant 5, mRl
 NM_175728 Homo sapiens interleukin 5 receptor, alpha (IL5RA), transcript variant 6, mRl
 NM_175729 Homo sapiens synovial sarcoma, X breakpoint 4 (SSX4), transcript variant 2,
 NM_175733 Homo sapiens synaptotagmin IX (SYT9), mRNA
 NM_175734 Homo sapiens hypothetical protein LOC201243 (LOC201243), mRNA
 NM_175735 Homo sapiens lysozyme-like (LYG2), mRNA
 NM_175736 Homo sapiens formin-like 3 (FMNL3), transcript variant 1, mRNA
 NM_175737 Homo sapiens klotho beta like (LOC152831), mRNA
 NM_175738 Homo sapiens RAB37, member RAS oncogene family (RAB37), mRNA
 NM_175739 Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1 anti
 NM_175741 Homo sapiens nuclear protein in testis (NUT), mRNA
 NM_175742 Homo sapiens melanoma antigen, family A, 2 (MAGEA2), transcript variant 2
 NM_175743 Homo sapiens melanoma antigen, family A, 2 (MAGEA2), transcript variant 3
 NM_175744 Homo sapiens ras homolog gene family, member C (RHOC), mRNA
 NM_175745 Homo sapiens cancer/testis antigen 3 (CTAG3), transcript variant 1, mRNA
 NM_175747 Homo sapiens oligodendrocyte transcription factor 3 (OLIG3), mRNA
 NM_175748 Homo sapiens chromosome 14 open reading frame 130 (C14orf130), mRNA
 NM_175767 Homo sapiens interleukin 6 signal transducer (gp130, oncostatin M receptor)
 NM_175768 Homo sapiens glutamate receptor, ionotropic, kainate 2 (GRIK2), transcript v

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NM_175834 Homo sapiens keratin 6L (KRT6L), mRNA
 NM_175839 Homo sapiens spermine oxidase (SMOX), transcript variant 1, mRNA
 NM_175840 Homo sapiens spermine oxidase (SMOX), transcript variant 2, mRNA
 NM_175841 Homo sapiens spermine oxidase (SMOX), transcript variant 3, mRNA
 NM_175842 Homo sapiens spermine oxidase (SMOX), transcript variant 4, mRNA
 NM_175847 Homo sapiens polypyrimidine tract binding protein 1 (PTBP1), transcript vari
 NM_175848 Homo sapiens DNA (cytosine-5)-methyltransferase 3 beta (DNMT3B), trans
 NM_175849 Homo sapiens DNA (cytosine-5)-methyltransferase 3 beta (DNMT3B), trans
 NM_175850 Homo sapiens DNA (cytosine-5)-methyltransferase 3 beta (DNMT3B), trans
 NM_175851 Homo sapiens a disintegrin-like and metalloprotease (repolysin type) with th
 NM_175852 Homo sapiens taxilin (DKFZp451J0118), mRNA
 NM_175854 Homo sapiens PABP1-dependent poly A-specific ribonuclease subunit PAN2
 NM_175857 Homo sapiens keratin associated protein 8-1 (KRTAP8-1), mRNA
 NM_175858 Homo sapiens keratin associated protein 11-1 (KRTAP11-1), mRNA
 NM_175859 Homo sapiens CTP synthase II (CTPS2), transcript variant 2, mRNA
 NM_175861 Homo sapiens ARG99 protein (ARG99), mRNA
 NM_175862 Homo sapiens CD86 antigen (CD28 antigen ligand 2, B7-2 antigen) (CD86),
 NM_175863 Homo sapiens AT rich interactive domain 1B (SWI1-like) (ARID1B), transcript
 NM_175864 Homo sapiens core-binding factor, runt domain, alpha subunit 2; translocate
 NM_175865 Homo sapiens ELYS transcription factor-like protein TMBS62 (ELYS), mRNA
 NM_175866 Homo sapiens kinase interacting with leukemia-associated gene (statmin) (
 NM_175867 Homo sapiens DNA (cytosine-5)-methyltransferase 3-like (DNMT3L), trans
 NM_175868 Homo sapiens melanoma antigen, family A, 6 (MAGEA6), transcript variant 2
 NM_175870 Homo sapiens hypothetical protein LOC90925 (LOC90925), mRNA
 NM_175871 Homo sapiens hypothetical protein FLJ35119 (FLJ35119), mRNA
 NM_175872 Homo sapiens FLJ38451 protein (FLJ38451), mRNA
 NM_175873 Homo sapiens hypothetical protein LOC134548 (LOC134548), mRNA
 NM_175874 Homo sapiens hypothetical protein MGC47869 (MGC47869), mRNA
 NM_175875 Homo sapiens sine oculis homeobox homolog 5 (Drosophila) (SIX5), mRNA
 NM_175876 Homo sapiens exocyst complex component 8 (EXOC8), mRNA
 NM_175877 Homo sapiens hypothetical protein MGC35023 (MGC35023), mRNA
 NM_175878 Homo sapiens hypothetical protein MGC57211 (MGC57211), mRNA
 NM_175881 Homo sapiens hypothetical protein MGC48986 (MGC48986), mRNA
 NM_175882 Homo sapiens intramembrane protease 5 (IMP5), mRNA
 NM_175884 Homo sapiens hypothetical protein FLJ38031 (FLJ38031), mRNA
 NM_175885 Homo sapiens hypothetical protein MGC33846 (MGC33846), mRNA
 NM_175886 Homo sapiens phosphoribosyl pyrophosphate synthetase 1-like 1 (PRPS1L1
 NM_175887 Homo sapiens hypothetical protein LOC222171 (LOC222171), mRNA
 NM_175892 Homo sapiens hypothetical protein FLJ37266 (FLJ37266), mRNA
 NM_175895 Homo sapiens hypothetical protein FLJ25590 (FLJ25590), mRNA
 NM_175898 Homo sapiens hypothetical protein LOC283687 (LOC283687), mRNA
 NM_175900 Homo sapiens hypothetical protein FLJ35681 (FLJ35681), mRNA
 NM_175901 Homo sapiens hypothetical protein LOC283932 (LOC283932), mRNA
 NM_175902 Homo sapiens hypothetical protein FLJ22222 (FLJ22222), mRNA
 NM_175903 Homo sapiens hypothetical protein LOC284033 (LOC284033), mRNA
 NM_175904 Homo sapiens hypothetical protein FLJ40121 (FLJ40121), mRNA
 NM_175906 Homo sapiens hypothetical protein MGC33608 (MGC33608), mRNA
 NM_175907 Homo sapiens zinc binding alcohol dehydrogenase, domain containing 2 (ZA
 NM_175908 Homo sapiens hypothetical protein LOC284296 (LOC284296), mRNA
 NM_175910 Homo sapiens zinc finger protein 493 (ZNF493), mRNA
 NM_175911 Homo sapiens hypothetical protein MGC40047 (MGC40047), mRNA
 NM_175913 Homo sapiens junctophilin 2 (JPH2), transcript variant 2, mRNA
 NM_175918 Homo sapiens hypothetical protein FLJ34443 (FLJ34443), mRNA
 NM_175920 Homo sapiens hypothetical protein FLJ39485 (FLJ39485), mRNA
 NM_175921 Homo sapiens hypothetical protein LOC285636 (LOC285636), mRNA
 NM_175922 Homo sapiens hypothetical protein MGC35308 (MGC35308), mRNA
 NM_175923 Homo sapiens hypothetical protein MGC42630 (MGC42630), mRNA

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NM_176876 Homo sapiens purinergic receptor P2Y, G-protein coupled, 12 (P2RY12), tra
 NM_176877 Homo sapiens InaD-like protein (INADL), transcript variant 2, mRNA
 NM_176878 Homo sapiens InaD-like protein (INADL), transcript variant 4, mRNA
 NM_176880 Homo sapiens TR4 orphan receptor associated protein TRA16 (TPRA16), mR
 NM_176881 Homo sapiens taste receptor, type 2, member 39 (TAS2R39), mRNA
 NM_176882 Homo sapiens taste receptor, type 2, member 40 (TAS2R40), mRNA
 NM_176883 Homo sapiens taste receptor, type 2, member 41 (TAS2R41), mRNA
 NM_176884 Homo sapiens taste receptor, type 2, member 43 (TAS2R43), mRNA
 NM_176885 Homo sapiens taste receptor, type 2, member 44 (TAS2R44), mRNA
 NM_176886 Homo sapiens taste receptor, type 2, member 45 (TAS2R45), mRNA
 NM_176887 Homo sapiens taste receptor, type 2, member 46 (TAS2R46), mRNA
 NM_176888 Homo sapiens taste receptor, type 2, member 48 (TAS2R48), mRNA
 NM_176889 Homo sapiens taste receptor, type 2, member 49 (TAS2R49), mRNA
 NM_176890 Homo sapiens taste receptor, type 2, member 50 (TAS2R50), mRNA
 NM_176891 Homo sapiens interferon epsilon 1 (IFNE1), mRNA
 NM_176894 Homo sapiens purinergic receptor P2Y, G-protein coupled, 13 (P2RY13), tra
 NM_176895 Homo sapiens phosphatidic acid phosphatase type 2A (PPAP2A), transcript
 NM_177398 Homo sapiens LIM homeobox transcription factor 1, alpha (LMX1A), transcr
 NM_177399 Homo sapiens LIM homeobox transcription factor 1, alpha (LMX1A), transcr
 NM_177400 Homo sapiens NK6 transcription factor related, locus 2 (Drosophila) (NKX6-2
 NM_177402 Homo sapiens synaptotagmin II (SYT2), mRNA
 NM_177403 Homo sapiens RAB7B, member RAS oncogene family (RAB7B), mRNA
 NM_177404 Homo sapiens melanoma antigen, family B, 1 (MAGEB1), transcript variant 2
 NM_177405 Homo sapiens cat eye syndrome chromosome region, candidate 1 (CECR1),
 NM_177414 Homo sapiens phosphatidic acid phosphatase type 2B (PPAP2B), transcript
 NM_177415 Homo sapiens melanoma antigen, family B, 1 (MAGEB1), transcript variant 3
 NM_177417 Homo sapiens kinesin light chain 2-like (KLC2L), transcript variant 1, mRNA
 NM_177422 Homo sapiens eukaryotic translation initiation factor 2C, 3 (EIF2C3), transcr
 NM_177423 Homo sapiens protein tyrosine phosphatase, receptor type, f polypeptide (PT
 NM_177424 Homo sapiens syntaxin 12 (STX12), mRNA
 NM_177427 Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 7 (P2RX7)
 NM_177433 Homo sapiens melanoma antigen, family D, 2 (MAGED2), transcript variant 2
 NM_177434 Homo sapiens FtsJ homolog 1 (E. coli) (FTSJ1), transcript variant 2, mRNA
 NM_177435 Homo sapiens peroxisome proliferative activated receptor, delta (PPARD), tr
 NM_177436 Homo sapiens CSE1 chromosome segregation 1-like (yeast) (CSE1L), trans
 NM_177437 Homo sapiens taste receptor, type 2, member 60 (TAS2R60), mRNA
 NM_177438 Homo sapiens Dicer1, Dcr-1 homolog (Drosophila) (DICER1), transcript varie
 NM_177439 Homo sapiens FtsJ homolog 1 (E. coli) (FTSJ1), transcript variant 3, mRNA
 NM_177441 Homo sapiens hypothetical protein MGC3123 (MGC3123), mRNA
 NM_177442 Homo sapiens FtsJ homolog 2 (E. coli) (FTSJ2), transcript variant 2, mRNA
 NM_177444 Homo sapiens PTPRF Interacting protein, binding protein 1 (liprin beta 1) (PF
 NM_177452 Homo sapiens trafficking protein particle complex 6B (TRAPPC6B), mRNA
 NM_177453 Homo sapiens progesterone and adiponectin receptor family member III (PAQR3), m
 NM_177454 Homo sapiens KIAA1946 (KIAA1946), mRNA
 NM_177455 Homo sapiens class II bHLH protein MIST1 (MIST1), mRNA
 NM_177456 Homo sapiens melanoma antigen, family C, 3 (MAGEC3), transcript variant 2
 NM_177457 Homo sapiens Ly-6 neurotoxin-like protein 1 (LYNX1), transcript variant 3, m
 NM_177458 Homo sapiens secreted Ly6/uPAR related protein 2 (SLURP2), mRNA
 NM_177476 Homo sapiens Ly-6 neurotoxin-like protein 1 (LYNX1), transcript variant 4, m
 NM_177477 Homo sapiens Ly-6 neurotoxin-like protein 1 (LYNX1), transcript variant 5, m
 NM_177478 Homo sapiens ferritin mitochondrial (FTMT), mRNA
 NM_177483 Homo sapiens glycosylphosphatidylinositol specific phospholipase D1 (GPLI
 NM_177524 Homo sapiens mesoderm specific transcript homolog (mouse) (MEST), trans
 NM_177525 Homo sapiens mesoderm specific transcript homolog (mouse) (MEST), trans
 NM_177526 Homo sapiens phosphatidic acid phosphatase type 2C (PPAP2C), transcript
 NM_177528 Homo sapiens sulfotransferase family, cytosolic, 1A, phenol-prefering, mem
 NM_177529 Homo sapiens sulfotransferase family, cytosolic, 1A, phenol-prefering, mem

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NM_177530 Homo sapiens sulfotransferase family, cytosolic, 1A, phenol-preferring, mem
 NM_177531 Homo sapiens polycystic kidney and hepatic disease 1 (autosomal recessive
 NM_177532 Homo sapiens Ras association (RalGDS/AF-6) domain family 6 (RASSF6), t
 NM_177533 Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 14
 NM_177534 Homo sapiens sulfotransferase family, cytosolic, 1A, phenol-preferring, mem
 NM_177535 Homo sapiens melanoma antigen, family D, 4 (MAGED4), transcript variant 2
 NM_177536 Homo sapiens sulfotransferase family, cytosolic, 1A, phenol-preferring, mem
 NM_177537 Homo sapiens melanoma antigen, family D, 4 (MAGED4), transcript variant 3
 NM_177538 Homo sapiens cytochrome P450, family 20, subfamily A, polypeptide 1 (CYP
 NM_177539 Homo sapiens taste receptor, type 1, member 1 (TAS1R1), transcript variant
 NM_177540 Homo sapiens taste receptor, type 1, member 1 (TAS1R1), transcript variant
 NM_177541 Homo sapiens taste receptor, type 1, member 1 (TAS1R1), transcript variant
 NM_177542 Homo sapiens small nuclear ribonucleoprotein D2 polypeptide 16.5kDa (SNF
 NM_177543 Homo sapiens phosphatidic acid phosphatase type 2C (PPAP2C), transcript
 NM_177549 Homo sapiens M8 protein (LOC149830), mRNA
 NM_177550 Homo sapiens solute carrier family 13 (sodium-dependent citrate transporter
 NM_177551 Homo sapiens G protein-coupled receptor 109A (GPR109A), mRNA
 NM_177552 Homo sapiens sulfotransferase family, cytosolic, 1A, phenol-preferring, mem
 NM_177553 Homo sapiens growth arrest-specific 2 (GAS2), transcript variant 2, mRNA
 NM_177554 Homo sapiens acid phosphatase 1, soluble (ACP1), transcript variant 1, mRNA
 NM_177555 Homo sapiens trophinin (TRO), transcript variant 1, mRNA
 NM_177556 Homo sapiens trophinin (TRO), transcript variant 2, mRNA
 NM_177557 Homo sapiens trophinin (TRO), transcript variant 5, mRNA
 NM_177558 Homo sapiens trophinin (TRO), transcript variant 4, mRNA
 NM_177559 Homo sapiens casein kinase 2, alpha 1 polypeptide (CSNK2A1), transcript v
 NM_177560 Homo sapiens casein kinase 2, alpha 1 polypeptide (CSNK2A1), transcript v
 NM_177924 Homo sapiens N-acylsphingosine amidohydrolase (acid ceramidase) 1 (ASA
 NM_177925 Homo sapiens H2A histone family, member J (H2AFJ), transcript variant 2, m
 NM_177926 Homo sapiens CSRP2 binding protein (CSRP2BP), transcript variant 2, mRNA
 NM_177937 Homo sapiens golgi phosphoprotein 2 (GOLPH2), transcript variant 2, mRNA
 NM_177938 Homo sapiens hypoxia-inducible factor prolyl 4-hydroxylase (PH-4), mRNA
 NM_177939 Homo sapiens hypoxia-inducible factor prolyl 4-hydroxylase (PH-4), transcript
 NM_177947 Homo sapiens armadillo repeat containing, X-linked 3 (ARMCX3), transcript
 NM_177948 Homo sapiens armadillo repeat containing, X-linked 3 (ARMCX3), transcript
 NM_177949 Homo sapiens armadillo repeat containing, X-linked 2 (ARMCX2), mRNA
 NM_177951 Homo sapiens protein phosphatase 1A (formerly 2C), magnesium-dependen
 NM_177952 Homo sapiens protein phosphatase 1A (formerly 2C), magnesium-dependen
 NM_177953 Homo sapiens dynein, cytoplasmic, light polypeptide 2A (DNCL2A), transcript
 NM_177954 Homo sapiens dynein, cytoplasmic, light polypeptide 2A (DNCL2A), transcript
 NM_177959 Homo sapiens dock protein 5 (DOK5), transcript variant 2, mRNA
 NM_177963 Homo sapiens synaptotagmin XII (SYT12), mRNA
 NM_177964 Homo sapiens hypothetical protein LOC130576 (LOC130576), mRNA
 NM_177965 Homo sapiens hypothetical protein LOC157857 (LOC157857), mRNA
 NM_177966 Homo sapiens hypothetical protein DKFZp667B1218 (DKFZp667B1218), mF
 NM_177967 Homo sapiens phosphoglycerate dehydrogenase like 1 (PHGDHL1), mRNA
 NM_177968 Homo sapiens protein phosphatase 1B (formerly 2C), magnesium-dependen
 NM_177969 Homo sapiens protein phosphatase 1B (formerly 2C), magnesium-dependen
 NM_177972 Homo sapiens tubby homolog (mouse) (TUB), transcript variant 2, mRNA
 NM_177973 Homo sapiens sulfotransferase family, cytosolic, 2B, member 1 (SULT2B1), l
 NM_177974 Homo sapiens H63 breast cancer expressed gene (H63), transcript variant 2
 NM_177976 Homo sapiens ADP-ribosylation factor-like 6 (ARL6), transcript variant 2, mR
 NM_177977 Homo sapiens huntingtin-associated protein 1 (neuroan 1) (HAP1), transcript
 NM_177978 Homo sapiens chordin (CHRD), transcript variant 2, mRNA
 NM_177979 Homo sapiens chordin (CHRD), transcript variant 3, mRNA
 NM_177980 Homo sapiens cadherin-like 26 (CDH26), transcript variant a, mRNA
 NM_177983 Homo sapiens protein phosphatase 1G (formerly 2C), magnesium-dependen
 NM_177985 Homo sapiens ADP-ribosylation factor-like 5 (ARL5), transcript variant 2, mR

NM_177986 Homo sapiens desmoglein 4 (DSG4), mRNA
 NM_177987 Homo sapiens tubulin, beta 8 (TUBB8-pending), mRNA
 NM_177988 Homo sapiens mitochondrial ribosomal protein L47 (MRPL47), nuclear gene
 NM_177989 Homo sapiens actin-like 6A (ACTL6A), transcript variant 2, mRNA
 NM_177990 Homo sapiens p21(CDKN1A)-activated kinase 7 (PAK7), transcript variant 2,
 NM_177991 Homo sapiens dual specificity phosphatase-like 15 (DUSP15), transcript vari
 NM_177995 Homo sapiens protein tyrosine phosphatase domain containing 1 (PTPDC1),
 NM_177996 Homo sapiens erythrocyte membrane protein band 4.1-like 1 (EPB41L1), tra
 NM_177998 Homo sapiens otopetrin 1 (OTOP1), mRNA
 NM_177999 Homo sapiens ankyrin repeat and SOCS box-containing 6 (ASB6), transcript
 NM_178000 Homo sapiens protein phosphatase 2A, regulatory subunit B' (PR 53) (PPP2
 NM_178001 Homo sapiens protein phosphatase 2A, regulatory subunit B' (PR 53) (PPP2
 NM_178002 Homo sapiens protein phosphatase 2A, regulatory subunit B' (PR 53) (PPP2
 NM_178003 Homo sapiens protein phosphatase 2A, regulatory subunit B' (PR 53) (PPP2
 NM_178004 Homo sapiens proline rich membrane anchor 1 (PRIMA1), transcript variant I
 NM_178006 Homo sapiens START domain containing 13 (STARD13), transcript variant a
 NM_178007 Homo sapiens START domain containing 13 (STARD13), transcript variant b
 NM_178008 Homo sapiens START domain containing 13 (STARD13), transcript variant d
 NM_178009 Homo sapiens diacylglycerol kinase, eta (DGKH), transcript variant 2, mRNA
 NM_178010 Homo sapiens SRY (sex determining region Y)-box 5 (SOX5), transcript vari
 NM_178011 Homo sapiens leucine rich repeat transmembrane neuronal 3 (LRRTM3), mF
 NM_178012 Homo sapiens tubulin, beta polypeptide paralog (MGC8685), mRNA
 NM_178013 Homo sapiens proline rich membrane anchor 1 (PRIMA1), mRNA
 NM_178014 Homo sapiens beta 5-tubulin (OK/SW-cl.56), mRNA
 NM_178019 Homo sapiens cation channel, sperm associated 3 (CATSPER3), mRNA
 NM_178025 Homo sapiens gamma-glutamyltransferase-like 3 (GGTL3), transcript variant
 NM_178028 Homo sapiens gamma-glutamyltransferase-like 3 (GGTL3), transcript variant
 NM_178031 Homo sapiens heat shock 70kDa protein 5 (glucose-regulated protein, 78kD;
 NM_178033 Homo sapiens cytochrome P450, family 4, subfamily X, polypeptide 1 (CYP4
 NM_178034 Homo sapiens phospholipase A2, group IVD (cytosolic) (PLA2G4D), mRNA
 NM_178037 Homo sapiens Rab6-interacting protein 2 (ELKS), transcript variant beta, mR
 NM_178038 Homo sapiens Rab6-interacting protein 2 (ELKS), transcript variant gamma,
 NM_178039 Homo sapiens Rab6-interacting protein 2 (ELKS), transcript variant delta, mF
 NM_178040 Homo sapiens Rab6-interacting protein 2 (ELKS), transcript variant epsilon, I
 NM_178042 Homo sapiens actin-like 6A (ACTL6A), transcript variant 3, mRNA
 NM_178043 Homo sapiens FLJ10378 protein (FLJ10378), transcript variant 2, mRNA
 NM_178044 Homo sapiens hypothetical protein MGC5179 (MGC5178), transcript variant
 NM_178120 Homo sapiens distal-less homeo box 1 (DLX1), mRNA
 NM_178121 Homo sapiens HBV pre-s2 binding protein 1 (SBP1), mRNA
 NM_178122 Homo sapiens hypothetical protein LOC90529 (LOC90529), mRNA
 NM_178123 Homo sapiens SEC14 and spectrin domains 1 (SESTD1), mRNA
 NM_178124 Homo sapiens hypothetical protein LOC91966 (LOC91966), mRNA
 NM_178125 Homo sapiens tripartite motif-containing 50A (TRIM50A), mRNA
 NM_178128 Homo sapiens hypothetical protein LOC162427 (LOC162427), mRNA
 NM_178127 Homo sapiens angiotensin-like 5 (ANGPTL5), mRNA
 NM_178128 Homo sapiens similar to delta 5 fatty acid desaturase (LOC283985), mRNA
 NM_178129 Homo sapiens purinergic receptor P2Y, G-protein coupled, 8 (P2RY8), mRN
 NM_178130 Homo sapiens thioredoxin domain containing 6 (TXNDC6), mRNA
 NM_178134 Homo sapiens cytochrome P450, family 4, subfamily Z, polypeptide 1 (CYP4
 NM_178135 Homo sapiens short-chain dehydrogenase/reductase 9 (SCDR9), mRNA
 NM_178136 Homo sapiens polymerase (DNA-directed), delta interacting protein 3 (POLD
 NM_178138 Homo sapiens LIM homeobox 3 (LHX3), transcript variant 1, mRNA
 NM_178140 Homo sapiens PDZ domain containing 3 (PDZK3), transcript variant 1, mRN
 NM_178145 Homo sapiens Ras association (RalGDS/AF-6) domain family 4 (RASSF4), t
 NM_178148 Homo sapiens solute carrier family 35, member B2 (SLC35B2), mRNA
 NM_178150 Homo sapiens F-box protein, helicase, 18 (FBXO18), transcript variant 2, mF
 NM_178151 Homo sapiens doublecortin; lissencephaly, X-linked (doublecortin) (DCX), tr

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NM_178152 Homo sapiens doublecortin; lissencephaly, X-linked (doublecortin) (DCX), tn
 NM_178153 Homo sapiens doublecortin; lissencephaly, X-linked (doublecortin) (DCX), tn
 NM_178154 Homo sapiens fucosyltransferase 8 (alpha (1,6) fucosyltransferase) (FUT8), l
 NM_178155 Homo sapiens fucosyltransferase 8 (alpha (1,6) fucosyltransferase) (FUT8), l
 NM_178156 Homo sapiens fucosyltransferase 8 (alpha (1,6) fucosyltransferase) (FUT8), l
 NM_178157 Homo sapiens fucosyltransferase 8 (alpha (1,6) fucosyltransferase) (FUT8), l
 NM_178159 Homo sapiens hypothetical protein FLJ12949 (FLJ12949), transcript variant :
 NM_178160 Homo sapiens otopetrin 2 (OTOP2), mRNA
 NM_178161 Homo sapiens pancreas specific transcription factor, 1a (PTF1A), mRNA
 NM_178167 Homo sapiens zinc finger protein 598 (ZNF598), mRNA
 NM_178168 Homo sapiens olfactory receptor, family 10, subfamily A, member 5 (OR10A)
 NM_178169 Homo sapiens Ras association (RalGDS/AF-6) domain family 3 (RASSF3), n
 NM_178170 Homo sapiens NIMA (never in mitosis gene a)-related kinase 8 (NEK8), mRi
 NM_178171 Homo sapiens gasdemlin 1 (GSDM1), mRNA
 NM_178172 Homo sapiens high density lipoprotein-binding protein (LOC338328), mRNA
 NM_178173 Homo sapiens hypothetical protein LOC339834 (LOC339834), mRNA
 NM_178174 Homo sapiens triggering receptor expressed on myeloid cells-like 1 (TREM)
 NM_178175 Homo sapiens lipoma HMGIC fusion partner-like 1 (LHFPL1), mRNA
 NM_178176 Homo sapiens monoacylglycerol O-acyltransferase 3 (MOGAT3), mRNA
 NM_178177 Homo sapiens nicotinamide nucleotide adenyltransferase 3 (NMNAT3), mF
 NM_178181 Homo sapiens CUB domain-containing protein 1 (CDCP1), transcript variant
 NM_178190 Homo sapiens ATPase inhibitory factor 1 (ATPIF1), nuclear gene encoding n
 NM_178191 Homo sapiens ATPase inhibitory factor 1 (ATPIF1), nuclear gene encoding n
 NM_178221 Homo sapiens APG4 autophagy 4 homolog C (S. cerevisiae) (APG4C), trans
 NM_178225 Homo sapiens F-box and WD-40 domain protein 5 (FBXW5), transcript varia
 NM_178226 Homo sapiens F-box and WD-40 domain protein 5 (FBXW5), transcript varia
 NM_178228 Homo sapiens killer cell immunoglobulin-like receptor, two domains, short C
 NM_178229 Homo sapiens IQ motif containing GTPase activating protein 3 (IQGAP3), mi
 NM_178230 Homo sapiens cyclophilin-LC (COAS2), mRNA
 NM_178231 Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region,
 NM_178232 Homo sapiens hyaluronan and proteoglycan link protein 3 (HAPLN3), mRNA
 NM_178233 Homo sapiens otopetrin 3 (OTOP3), mRNA
 NM_178234 Homo sapiens tumor suppressor candidate 3 (TUSC3), transcript variant 2, n
 NM_178237 Homo sapiens SEC3-like 1 (S. cerevisiae) (SEC3L1), transcript variant 2, mF
 NM_178238 Homo sapiens paired immunoglobulin-like type 2 receptor beta (PILRB), trans
 NM_178270 Homo sapiens APG4 autophagy 4 homolog A (S. cerevisiae) (APG4A), trans
 NM_178271 Homo sapiens APG4 autophagy 4 homolog A (S. cerevisiae) (APG4A), trans
 NM_178272 Homo sapiens paired immunoglobulin-like type 2 receptor alpha (PILRA), trans
 NM_178273 Homo sapiens paired immunoglobulin-like type 2 receptor alpha (PILRA), trans
 NM_178275 Homo sapiens eEF1A2 binding protein (DKFZp434B1231), mRNA
 NM_178276 Homo sapiens chromosome 5 open reading frame 12 (C5orf12), mRNA
 NM_178311 Homo sapiens gamma-glutamyltransferase-like activity 4 (GGTLA4), transcr
 NM_178312 Homo sapiens gamma-glutamyltransferase-like activity 4 (GGTLA4), transcr
 NM_178313 Homo sapiens spectrin, beta, non-erythrocytic 1 (SPTBN1), transcript variant
 NM_178314 Homo sapiens hypothetical protein FLJ39378 (FLJ39378), mRNA
 NM_178324 Homo sapiens serine palmitoyltransferase, long chain base subunit 1 (SPTL)
 NM_178326 Homo sapiens APG4 autophagy 4 homolog B (S. cerevisiae) (APG4B), trans
 NM_178329 Homo sapiens chemokine (C-C motif) receptor 3 (CCR3), transcript variant 2
 NM_178331 Homo sapiens gonadotropin-releasing hormone 2 (GNRH2), transcript variat
 NM_178332 Homo sapiens gonadotropin-releasing hormone 2 (GNRH2), transcript variat
 NM_178335 Homo sapiens chromosome 3 open reading frame 6 (C3orf6), transcript variat
 NM_178336 Homo sapiens mitochondrial ribosomal protein L52 (MRPL52), nuclear gene
 NM_178338 Homo sapiens AP20 region protein (APRG1), transcript variant A, mRNA
 NM_178339 Homo sapiens AP20 region protein (APRG1), transcript variant B, mRNA
 NM_178340 Homo sapiens AP20 region protein (APRG1), transcript variant C, mRNA
 NM_178341 Homo sapiens AP20 region protein (APRG1), transcript variant D, mRNA
 NM_178342 Homo sapiens AP20 region protein (APRG1), transcript variant E, mRNA

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NM_178343 Homo sapiens AP20 region protein (APRG1), transcript variant F, mRNA
 NM_178344 Homo sapiens AP20 region protein (APRG1), transcript variant G, mRNA
 NM_178348 Homo sapiens late envelope protein 1 (LEP1), mRNA
 NM_178349 Homo sapiens small proline rich-like (epidermal differentiation complex) 2A (LOC178349), mRNA
 NM_178351 Homo sapiens late envelope protein 3 (LEP3), mRNA
 NM_178352 Homo sapiens late envelope protein 4 (LEP4), mRNA
 NM_178353 Homo sapiens late envelope protein 5 (LEP5), mRNA
 NM_178354 Homo sapiens late envelope protein 6 (LEP6), mRNA
 NM_178356 Homo sapiens small proline rich-like (epidermal differentiation complex) 4A (LOC178356), mRNA
 NM_178422 Homo sapiens membrane progesterin receptor alpha (MPRA), mRNA
 NM_178423 Homo sapiens histone deacetylase 9 (HDAC9), transcript variant 4, mRNA
 NM_178424 Homo sapiens SRY (sex determining region Y)-box 30 (SOX30), transcript variant 1, mRNA
 NM_178425 Homo sapiens histone deacetylase 9 (HDAC9), transcript variant 5, mRNA
 NM_178426 Homo sapiens aryl hydrocarbon receptor nuclear translocator (ARNT), transcript variant 1, mRNA
 NM_178427 Homo sapiens aryl hydrocarbon receptor nuclear translocator (ARNT), transcript variant 2, mRNA
 NM_178428 Homo sapiens late envelope protein 9 (LEP9), mRNA
 NM_178429 Homo sapiens late envelope protein 11 (LEP11), mRNA
 NM_178430 Homo sapiens small proline rich-like (epidermal differentiation complex) 1A (LOC178430), mRNA
 NM_178431 Homo sapiens late envelope protein 13 (LEP13), mRNA
 NM_178432 Homo sapiens cell cycle related kinase (CCRK), transcript variant 1, mRNA
 NM_178433 Homo sapiens late envelope protein 14 (LEP14), mRNA
 NM_178434 Homo sapiens small proline rich-like (epidermal differentiation complex) 3A (LOC178434), mRNA
 NM_178435 Homo sapiens late envelope protein 17 (LEP17), mRNA
 NM_178438 Homo sapiens small proline rich-like (epidermal differentiation complex) 5A (LOC178438), mRNA
 NM_178439 Homo sapiens germ cell-less homolog 1 (Drosophila) (GCL), mRNA
 NM_178441 Homo sapiens zinc finger, FYVE domain containing 1 (ZFVE1), transcript variant 1, mRNA
 NM_178443 Homo sapiens UNC-112 related protein 2 (URP2), transcript variant URP2LF, mRNA
 NM_178445 Homo sapiens chemokine (C-C motif) receptor-like 1 (CCR1), transcript variant 1, mRNA
 NM_178448 Homo sapiens chromosome 9 open reading frame 140 (C9orf140), mRNA
 NM_178449 Homo sapiens tuberin/fundinbular 39 residue protein precursor (TIP39), mRNA
 NM_178450 Homo sapiens hypothetical protein MGC48332 (MGC48332), mRNA
 NM_178451 Homo sapiens zinc finger, MYND domain containing 17 (ZMYND17), mRNA
 NM_178452 Homo sapiens similar to RIKEN cDNA 4930457P18 (LOC123872), mRNA
 NM_178453 Homo sapiens hypothetical protein MGC52282 (MGC52282), mRNA
 NM_178454 Homo sapiens hypothetical protein MGC54289 (MGC54289), mRNA
 NM_178456 Homo sapiens chromosome 20 open reading frame 85 (C20orf85), mRNA
 NM_178460 Homo sapiens protein tyrosine phosphatase, non-receptor type substrate 1-ii (LOC178460), mRNA
 NM_178463 Homo sapiens chromosome 20 open reading frame 166 (C20orf166), mRNA
 NM_178465 Homo sapiens TSPY-like 3 (TSPYL3), mRNA
 NM_178466 Homo sapiens chromosome 20 open reading frame 71 (C20orf71), mRNA
 NM_178467 Homo sapiens high-mobility group (nonhistone chromosomal) protein 4-like (LOC178467), mRNA
 NM_178468 Homo sapiens chromosome 20 open reading frame 128 (C20orf128), mRNA
 NM_178470 Homo sapiens WD repeat domain 40B (WDR40B), mRNA
 NM_178471 Homo sapiens G protein-coupled receptor 119 (GPR119), mRNA
 NM_178472 Homo sapiens chromosome 20 open reading frame 53 (C20orf53), mRNA
 NM_178477 Homo sapiens chromosome 20 open reading frame 179 (C20orf179), mRNA
 NM_178483 Homo sapiens chromosome 20 open reading frame 79 (C20orf79), mRNA
 NM_178491 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing-1 (LOC178491), mRNA
 NM_178493 Homo sapiens hypothetical protein LOC147111 (LOC147111), mRNA
 NM_178494 Homo sapiens hypothetical protein FLJ40125 (FLJ40125), mRNA
 NM_178495 Homo sapiens KIAA1754-like (KIAA1754), mRNA
 NM_178496 Homo sapiens similar to BcDNA:GH11415 gene product (LOC151963), mRNA
 NM_178497 Homo sapiens hypothetical protein FLJ23657 (FLJ23657), mRNA
 NM_178498 Homo sapiens hypothetical protein MGC52019 (MGC52019), mRNA
 NM_178499 Homo sapiens hypothetical protein MGC39827 (MGC39827), mRNA
 NM_178500 Homo sapiens phosphatase, orphan 1 (PHOSPHO1), mRNA
 NM_178502 Homo sapiens deltex 3 homolog (Drosophila) (DTX3), mRNA

NM_178504 Homo sapiens hypothetical protein FLJ40427 (FLJ40427), mRNA
 NM_178505 Homo sapiens transmembrane protein 26 (TMEM26), mRNA
 NM_178507 Homo sapiens NS5ATP13P2 protein (NS5ATP13P2), mRNA
 NM_178508 Homo sapiens hypothetical protein MGC57858 (MGC57858), mRNA
 NM_178509 Homo sapiens syntaxin binding protein 4 (STXBP4), mRNA
 NM_178510 Homo sapiens ankyrin repeat and kinase domain containing 1 (ANKK1), mRl
 NM_178514 Homo sapiens hypothetical protein LOC283487 (LOC283487), mRNA
 NM_178516 Homo sapiens hypothetical protein LOC283849 (LOC283849), mRNA
 NM_178517 Homo sapiens phosphatidylinositol glycan, class W (PIGW), mRNA
 NM_178518 Homo sapiens hypothetical protein FLJ36878 (FLJ36878), mRNA
 NM_178519 Homo sapiens hypothetical protein FLJ39421 (FLJ39421), mRNA
 NM_178520 Homo sapiens hypothetical protein FLJ38792 (FLJ38792), mRNA
 NM_178523 Homo sapiens zinc finger protein 616 (ZNF616), mRNA
 NM_178525 Homo sapiens hypothetical protein MGC33407 (MGC33407), mRNA
 NM_178527 Homo sapiens hypothetical protein MGC43026 (MGC43026), mRNA
 NM_178530 Homo sapiens hypothetical protein FLJ38379 (FLJ38379), mRNA
 NM_178532 Homo sapiens hypothetical protein LOC285671 (LOC285671), mRNA
 NM_178536 Homo sapiens lipocalin 12 (LCN12), mRNA
 NM_178537 Homo sapiens beta1,4-N-acetylgalactosaminyltransferases IV (Beta4GalNAc
 NM_178538 Homo sapiens hypothetical protein LOC338799 (LOC338799), mRNA
 NM_178539 Homo sapiens TAF42 protein (TAF42), mRNA
 NM_178540 Homo sapiens hypothetical protein MGC48915 (MGC48915), mRNA
 NM_178542 Homo sapiens hypothetical protein DKFZp782C2414 (DKFZp782C2414), mF
 NM_178543 Homo sapiens ecdonucleotide pyrophosphatase/phosphodiesterase 7 (ENPP
 NM_178544 Homo sapiens zinc finger protein 546 (ZNF546), mRNA
 NM_178545 Homo sapiens hypothetical protein LOC339456 (LOC339456), mRNA
 NM_178546 Homo sapiens hypothetical protein LOC339483 (LOC339483), mRNA
 NM_178547 Homo sapiens archease (ARCH), mRNA
 NM_178548 Homo sapiens adaptor-related protein complex 2, epsilon subunit (AP2E), ml
 NM_178549 Homo sapiens hypothetical protein MGC42493 (MGC42493), mRNA
 NM_178550 Homo sapiens hypothetical protein MGC48998 (MGC48998), mRNA
 NM_178552 Homo sapiens hypothetical protein MGC35206 (MGC35206), mRNA
 NM_178553 Homo sapiens hypothetical protein MGC44505 (MGC44505), mRNA
 NM_178554 Homo sapiens kyphoscoliosis peptidase (KY), mRNA
 NM_178555 Homo sapiens hypothetical protein FLJ25770 (FLJ25770), mRNA
 NM_178556 Homo sapiens hypothetical protein FLJ36180 (FLJ36180), mRNA
 NM_178557 Homo sapiens hypothetical protein FLJ37478 (FLJ37478), mRNA
 NM_178558 Homo sapiens hypothetical protein FLJ90430 (FLJ90430), mRNA
 NM_178559 Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 5 (A
 NM_178562 Homo sapiens hypothetical protein MGC50844 (MGC50844), mRNA
 NM_178563 Homo sapiens hypothetical protein LOC340351 (LOC340351), mRNA
 NM_178564 Homo sapiens hypothetical protein LOC340371 (LOC340371), mRNA
 NM_178565 Homo sapiens hypothetical protein MGC35555 (MGC35555), mRNA
 NM_178566 Homo sapiens zinc finger, DHHC domain containing 21 (ZDHHC21), mRNA
 NM_178568 Homo sapiens reticulum 4 receptor-like 1 (RTN4RL1), mRNA
 NM_178569 Homo sapiens CE1 protein (CE1), mRNA
 NM_178570 Homo sapiens reticulum 4 receptor-like 2 (RTN4RL2), mRNA
 NM_178571 Homo sapiens hypothetical protein MGC51025 (MGC51025), mRNA
 NM_178578 Homo sapiens proteasome (prosome, macropain) inhibitor subunit 1 (Pi31) (
 NM_178579 Homo sapiens proteasome (prosome, macropain) inhibitor subunit 1 (Pi31) (
 NM_178580 Homo sapiens histocompatibility (minor) 13 (HM13), transcript variant 2, mRl
 NM_178581 Homo sapiens histocompatibility (minor) 13 (HM13), transcript variant 3, mRl
 NM_178582 Homo sapiens histocompatibility (minor) 13 (HM13), transcript variant 4, mRl
 NM_178583 Homo sapiens WD repeat and FYVE domain containing 3 (WDFY3), transcr
 NM_178584 Homo sapiens septin 10 (SEPT10), transcript variant 2, mRNA
 NM_178585 Homo sapiens WD repeat and FYVE domain containing 3 (WDFY3), transcr
 NM_178586 Homo sapiens protein phosphatase 2, regulatory subunit B (B56), gamma iso

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NM_178587 Homo sapiens protein phosphatase 2, regulatory subunit B (B56), gamma iso
NM_178588 Homo sapiens protein phosphatase 2, regulatory subunit B (B56), gamma iso
NM_178812 Homo sapiens LYRIC/3D3 (LYRIC), mRNA
NM_178813 Homo sapiens A-kinase anchoring protein 28 (AKAP28), mRNA
NM_178814 Homo sapiens adaptor-related protein complex 1, sigma 3 subunit (AP1S3),
NM_178815 Homo sapiens ADP-ribosylation factor-like 8 (ARL8), mRNA
NM_178816 Homo sapiens cancer susceptibility candidate 2 (CASC2), mRNA
NM_178817 Homo sapiens chromosome 21 open reading frame 61 (C21orf61), transcript
NM_178818 Homo sapiens chemokine-like factor super family 4 (CKLFSF4), transcript va
NM_178819 Homo sapiens putative lysophosphatidic acid acyltransferase (DKFZp586M1
NM_178820 Homo sapiens F-box protein 27 (FBXO27), mRNA
NM_178821 Homo sapiens hypothetical protein FLJ25955 (FLJ25955), mRNA
NM_178822 Homo sapiens immunoglobulin superfamily, member 10 (IGSF10), mRNA
NM_178823 Homo sapiens chromosome 6 open reading frame 165 (C6orf165), mRNA
NM_178824 Homo sapiens hypothetical protein FLJ33620 (FLJ33620), mRNA
NM_178826 Homo sapiens transmembrane protein 16D (TMEM16D), mRNA
NM_178827 Homo sapiens hypothetical protein FLJ35834 (FLJ35834), mRNA
NM_178828 Homo sapiens chromosome 9 open reading frame 79 (C9orf79), mRNA
NM_178829 Homo sapiens chromosome 7 open reading frame 34 (C7orf34), mRNA
NM_178830 Homo sapiens hypothetical protein FLJ36888 (FLJ36888), mRNA
NM_178831 Homo sapiens opposite strand transcription unit to STAG3 (GATS), mRNA
NM_178832 Homo sapiens chromosome 10 open reading frame 83 (C10orf83), mRNA
NM_178833 Homo sapiens hypothetical protein BC009732 (LOC133308), mRNA
NM_178834 Homo sapiens laylin (LOC143903), mRNA
NM_178835 Homo sapiens hypothetical protein LOC152485 (LOC152485), mRNA
NM_178836 Homo sapiens similar to CG12314 gene product (LOC201164), mRNA
NM_178837 Homo sapiens similar to hypothetical testis protein from macaque (LOC3529
NM_178838 Homo sapiens hypothetical protein LOC90768 (MGC45800), mRNA
NM_178839 Homo sapiens leucine rich repeat transmembrane neuronal 1 (LRRTM1), mF
NM_178840 Homo sapiens hypothetical protein MGC24047 (MGC24047), mRNA
NM_178841 Homo sapiens ring finger protein 166 (RNF166), mRNA
NM_178842 Homo sapiens LAG1 longevity assurance homolog 3 (S. cerevisiae) (LASS3)
NM_178844 Homo sapiens NOD3 protein (NOD3), mRNA
NM_178849 Homo sapiens hepatocyte nuclear factor 4, alpha (HNF4A), transcript variant
NM_178850 Homo sapiens hepatocyte nuclear factor 4, alpha (HNF4A), transcript variant
NM_178857 Homo sapiens retinitis pigmentosa 1-like 1 (RP1L1), mRNA
NM_178858 Homo sapiens sideroflexin 2 (SFXN2), mRNA
NM_178859 Homo sapiens organic solute transporter beta (OSTbeta), mRNA
NM_178860 Homo sapiens seizure related 8 homolog (mouse) (SEZ8), mRNA
NM_178861 Homo sapiens zinc finger protein 183-like 1 (ZNF183L1), mRNA
NM_178862 Homo sapiens source of Immunodominant MHC-associated peptides (SIMP)
NM_178863 Homo sapiens potassium channel tetramerisation domain containing 13 (KC
NM_178864 Homo sapiens HLH-PAS transcription factor NXF (NXF), mRNA
NM_178865 Homo sapiens tumor differentially expressed 2-like (TDE2L), mRNA
NM_178867 Homo sapiens sideroflexin 4 (SFXN4), transcript variant 2, mRNA
NM_178868 Homo sapiens chemokine-like factor super family 8 (CKLFSF8), mRNA
NM_180699 Homo sapiens U11/U12 snRNP 35K (U1SNRNPBP), transcript variant 3, mR
NM_180703 Homo sapiens U11/U12 snRNP 35K (U1SNRNPBP), transcript variant 4, mR
NM_180976 Homo sapiens protein phosphatase 2, regulatory subunit B (B56), delta isofo
NM_180977 Homo sapiens protein phosphatase 2, regulatory subunit B (B56), delta isofo
NM_180981 Homo sapiens mitochondrial ribosomal protein L52 (MRPL52), nuclear gene
NM_180982 Homo sapiens mitochondrial ribosomal protein L52 (MRPL52), nuclear gene
NM_180989 Homo sapiens intimin thickness-related receptor (ITR), mRNA
NM_180990 Homo sapiens ligand-gated ion channel subunit (LGIC2), mRNA
NM_180991 Homo sapiens solute carrier organic anion transporter family, member 4C1 (C
NM_181041 Homo sapiens polybromo 1 (PB1), mRNA
NM_181042 Homo sapiens polybromo 1 (PB1), mRNA

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NM_181050 Homo sapiens axin 1 (AXIN1), transcript variant 2, mRNA
 NM_181054 Homo sapiens hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix)
 NM_181076 Homo sapiens golgin-67 (GOLGIN-67), transcript variant 2, mRNA
 NM_181077 Homo sapiens golgin-67 (GOLGIN-67), transcript variant 3, mRNA
 NM_181078 Homo sapiens interleukin 21 receptor (IL21R), transcript variant 2, mRNA
 NM_181079 Homo sapiens interleukin 21 receptor (IL21R), transcript variant 3, mRNA
 NM_181093 Homo sapiens ezrin-binding partner PACE-1 (PACE-1), transcript variant 2, r
 NM_181265 Homo sapiens WD repeat domain 17 (WDR17), transcript variant 2, mRNA
 NM_181268 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181269 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181270 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181271 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181272 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181283 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181285 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181286 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181287 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181288 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181289 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181290 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181291 Homo sapiens WD repeat domain 20 (WDR20), transcript variant 1, mRNA
 NM_181292 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181293 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181294 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181295 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181296 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181297 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181298 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181299 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181300 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181301 Homo sapiens chemokine-like factor super family 1 (CKLFSF1), transcript va
 NM_181302 Homo sapiens WD repeat domain 20 (WDR20), transcript variant 4, mRNA
 NM_181304 Homo sapiens mitochondrial ribosomal protein L52 (MRPL52), nuclear gene
 NM_181305 Homo sapiens mitochondrial ribosomal protein L52 (MRPL52), nuclear gene
 NM_181306 Homo sapiens mitochondrial ribosomal protein L52 (MRPL52), nuclear gene
 NM_181307 Homo sapiens mitochondrial ribosomal protein L52 (MRPL52), nuclear gene
 NM_181308 Homo sapiens WD repeat domain 20 (WDR20), transcript variant 3, mRNA
 NM_181309 Homo sapiens interleukin 22 receptor, alpha 2 (IL22RA2), transcript variant 2
 NM_181310 Homo sapiens interleukin 22 receptor, alpha 2 (IL22RA2), transcript variant 3
 NM_181311 Homo sapiens tafazzin (cardiomyopathy, dilated 3A (X-linked); endocardial fi
 NM_181312 Homo sapiens tafazzin (cardiomyopathy, dilated 3A (X-linked); endocardial fi
 NM_181313 Homo sapiens tafazzin (cardiomyopathy, dilated 3A (X-linked); endocardial fi
 NM_181314 Homo sapiens tafazzin (cardiomyopathy, dilated 3A (X-linked); endocardial fi
 NM_181332 Homo sapiens neuroligin 4, X-linked (NLGN4X), transcript variant 2, mRNA
 NM_181333 Homo sapiens Rho GTPase activating protein 8 (ARHGAP8), transcript varia
 NM_181334 Homo sapiens Rho GTPase activating protein 8 (ARHGAP8), transcript varia
 NM_181335 Homo sapiens Rho GTPase activating protein 8 (ARHGAP8), transcript varia
 NM_181336 Homo sapiens LEM domain containing 2 (LEMD2), mRNA
 NM_181337 Homo sapiens kidney associated antigen 1 (KAAG1), mRNA
 NM_181339 Homo sapiens interleukin 24 (IL24), transcript variant 2, mRNA
 NM_181340 Homo sapiens WD repeat domain 21 (WDR21), transcript variant 2, mRNA
 NM_181341 Homo sapiens WD repeat domain 21 (WDR21), transcript variant 3, mRNA
 NM_181342 Homo sapiens FK506 binding protein 7 (FKBP7), transcript variant 2, mRNA
 NM_181349 Homo sapiens SMAD specific E3 ubiquitin protein ligase 1 (SMURF1), transc
 NM_181351 Homo sapiens neural cell adhesion molecule 1 (NCAM1), mRNA
 NM_181353 Homo sapiens inhibitor of DNA binding 1, dominant negative helix-loop-helix
 NM_181354 Homo sapiens oxidation resistance 1 (OXR1), mRNA

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NM_181510 Homo sapiens WAP four-disulfide core domain 8 (WFDC8), transcript variant
 NM_181512 Homo sapiens mitochondrial ribosomal protein L21 (MRPL21), nuclear gene
 NM_181513 Homo sapiens mitochondrial ribosomal protein L21 (MRPL21), nuclear gene
 NM_181514 Homo sapiens mitochondrial ribosomal protein L21 (MRPL21), nuclear gene
 NM_181515 Homo sapiens mitochondrial ribosomal protein L21 (MRPL21), nuclear gene
 NM_181519 Homo sapiens synaptotagmin XV (SYT15), transcript variant b, mRNA
 NM_181521 Homo sapiens chemokine-like factor super family 4 (CKLFSF4), transcript variant
 NM_181522 Homo sapiens WAP four-disulfide core domain 3 (WFDC3), transcript variant
 NM_181523 Homo sapiens phosphoinositide-3-kinase, regulatory subunit, polypeptide 1 (NM_181524
 Homo sapiens phosphoinositide-3-kinase, regulatory subunit, polypeptide 1 (NM_181525
 Homo sapiens WAP four-disulfide core domain 3 (WFDC3), transcript variant
 NM_181526 Homo sapiens myosin, light polypeptide 9, regulatory (MYL9), transcript variant
 NM_181527 Homo sapiens N-acetyltransferase 5 (ARD1 homolog, S. cerevisiae) (NAT5),
 NM_181528 Homo sapiens N-acetyltransferase 5 (ARD1 homolog, S. cerevisiae) (NAT5),
 NM_181530 Homo sapiens WAP four-disulfide core domain 3 (WFDC3), transcript variant
 NM_181531 Homo sapiens butyrophilin, subfamily 2, member A2 (BTN2A2), transcript variant
 NM_181532 Homo sapiens ES cell expressed Ras (ERAS), mRNA
 NM_181533 Homo sapiens chromosome 14 open reading frame 29 (C14orf29), transcript
 NM_181534 Homo sapiens keratin 25A (KRT25A), mRNA
 NM_181535 Homo sapiens keratin 25D (KRT25D), mRNA
 NM_181536 Homo sapiens polycystic kidney disease 1-like 3 (PKD1L3), mRNA
 NM_181537 Homo sapiens keratin 25C (KRT25C), mRNA
 NM_181538 Homo sapiens gap junction protein, epsilon 1, 29kDa (GJ1), mRNA
 NM_181539 Homo sapiens keratin 25B (KRT25B), mRNA
 NM_181552 Homo sapiens cut-1-like 1, CCAAT displacement protein (Drosophila) (CUTL1)
 NM_181553 Homo sapiens chemokine-like factor super family 3 (CKLFSF3), transcript variant
 NM_181554 Homo sapiens chemokine-like factor super family 3 (CKLFSF3), transcript variant
 NM_181555 Homo sapiens chemokine-like factor super family 3 (CKLFSF3), transcript variant
 NM_181558 Homo sapiens replication factor C (activator 1) 3, 38kDa (RFC3), transcript variant
 NM_181571 Homo sapiens cAMP responsive element modulator (CREM), transcript variant
 NM_181572 Homo sapiens regulator of G-protein signalling like 1 (RGS1), mRNA
 NM_181573 Homo sapiens replication factor C (activator 1) 4, 37kDa (RFC4), transcript variant
 NM_181575 Homo sapiens ancient ubiquitous protein 1 (AUP1), transcript variant 2, mRNA
 NM_181576 Homo sapiens ancient ubiquitous protein 1 (AUP1), transcript variant 3, mRNA
 NM_181578 Homo sapiens replication factor C (activator 1) 5, 36.5kDa (RFC5), transcript
 NM_181581 Homo sapiens protein similar to E.coli yhdg and R. capsulatus nifR3 (PP35),
 NM_181597 Homo sapiens uridine phosphorylase 1 (UPP1), transcript variant 2, mRNA
 NM_181598 Homo sapiens spastic paraplegia 3A (autosomal dominant) (SPG3A), mRNA
 NM_181599 Homo sapiens keratin associated protein 13-1 (KRTAP13-1), mRNA
 NM_181600 Homo sapiens keratin associated protein 13-4 (KRTAP13-4), mRNA
 NM_181602 Homo sapiens keratin associated protein 6-1 (KRTAP6-1), mRNA
 NM_181604 Homo sapiens keratin associated protein 6-2 (KRTAP6-2), mRNA
 NM_181605 Homo sapiens keratin associated protein 6-3 (KRTAP6-3), mRNA
 NM_181607 Homo sapiens keratin associated protein 19-1 (KRTAP19-1), mRNA
 NM_181608 Homo sapiens keratin associated protein 19-2 (KRTAP19-2), mRNA
 NM_181609 Homo sapiens keratin associated protein 19-3 (KRTAP19-3), mRNA
 NM_181610 Homo sapiens keratin associated protein 19-4 (KRTAP19-4), mRNA
 NM_181611 Homo sapiens keratin associated protein 19-5 (KRTAP19-5), mRNA
 NM_181612 Homo sapiens keratin associated protein 19-6 (KRTAP19-6), mRNA
 NM_181614 Homo sapiens keratin associated protein 19-7 (KRTAP19-7), mRNA
 NM_181615 Homo sapiens keratin associated protein 20-1 (KRTAP20-1), mRNA
 NM_181616 Homo sapiens keratin associated protein 20-2 (KRTAP20-2), mRNA
 NM_181617 Homo sapiens keratin associated protein 21-2 (KRTAP21-2), mRNA
 NM_181618 Homo sapiens chemokine-like factor super family 5 (CKLFSF5), transcript variant
 NM_181619 Homo sapiens keratin associated protein 21-1 (KRTAP21-1), mRNA
 NM_181620 Homo sapiens keratin associated protein 22-1 (KRTAP22-1), mRNA
 NM_181621 Homo sapiens keratin associated protein 13-2 (KRTAP13-2), nuclear gene e

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NM_181622 Homo sapiens keratin associated protein 13-3 (KRTAP13-3), mRNA
 NM_181623 Homo sapiens keratin associated protein 15-1 (KRTAP15-1), mRNA
 NM_181624 Homo sapiens keratin associated protein 23-1 (KRTAP23-1), mRNA
 NM_181640 Homo sapiens chemokine-like factor (CKLF), transcript variant 2, mRNA
 NM_181641 Homo sapiens chemokine-like factor (CKLF), transcript variant 4, mRNA
 NM_181642 Homo sapiens serine protease inhibitor, Kunitz type 1 (SPINT1), transcript va
 NM_181643 Homo sapiens hypothetical protein LOC128344 (LOC128344), mRNA
 NM_181644 Homo sapiens hypothetical protein DKFZp761N1114 (DKFZp761N1114), mF
 NM_181645 Homo sapiens hypothetical protein FLJ25393 (FLJ25393), mRNA
 NM_181646 Homo sapiens hypothetical protein FLJ32110 (FLJ32110), mRNA
 NM_181647 Homo sapiens hypothetical protein LOC285398 (LOC285398), mRNA
 NM_181651 Homo sapiens peroxiredoxin 5 (PRDX5), nuclear gene encoding mitochondri
 NM_181652 Homo sapiens peroxiredoxin 5 (PRDX5), nuclear gene encoding mitochondri
 NM_181654 Homo sapiens complexin 4 (CPLX4), mRNA
 NM_181655 Homo sapiens hypothetical protein LOC284018 (LOC284018), transcript vari
 NM_181656 Homo sapiens hypothetical protein LOC284018 (LOC284018), transcript vari
 NM_181657 Homo sapiens leukotriene B4 receptor (LTB4R), mRNA
 NM_181659 Homo sapiens nuclear receptor coactivator 3 (NCOA3), transcript variant 1, r
 NM_181661 Homo sapiens Cohen syndrome 1 (COH1), transcript variant 4, mRNA
 NM_181670 Homo sapiens E2a-Pbx1-associated protein (EB-1), transcript variant 2, mR
 NM_181671 Homo sapiens phosphatidylinositol transfer protein, cytoplasmic 1 (PITPNC1
 NM_181672 Homo sapiens O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-
 NM_181673 Homo sapiens O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-
 NM_181674 Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit B (PF
 NM_181675 Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit B (PF
 NM_181676 Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit B (PF
 NM_181677 Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit B (PF
 NM_181678 Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit B (PF
 NM_181679 Homo sapiens NFS1 nitrogen fixation 1 (S. cerevisiae) (NFS1), nuclear gene
 NM_181684 Homo sapiens keratin associated protein 12-2 (KRTAP12-2), mRNA
 NM_181686 Homo sapiens keratin associated protein 12-1 (KRTAP12-1), mRNA
 NM_181688 Homo sapiens keratin associated protein 10-10 (KRTAP10-10), mRNA
 NM_181689 Homo sapiens neuronatin (NNAT), transcript variant 2, mRNA
 NM_181690 Homo sapiens v-akt murine thymoma viral oncogene homolog 3 (protein kin
 NM_181698 Homo sapiens peroxiredoxin 1 (PRDX1), transcript variant 2, mRNA
 NM_181697 Homo sapiens peroxiredoxin 1 (PRDX1), transcript variant 3, mRNA
 NM_181698 Homo sapiens chromosome 10 open reading frame 9 (C10orf9), mRNA
 NM_181699 Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit A (PF
 NM_181701 Homo sapiens quiescin Q6-like 1 (QSCN6L1), mRNA
 NM_181702 Homo sapiens GTP binding protein overexpressed in skeletal muscle (GEM).
 NM_181703 Homo sapiens gap junction protein, alpha 5, 40kDa (connexin 40) (GJA5), tr
 NM_181704 Homo sapiens B melanoma antigen family, member 4 (BAGE4), mRNA
 NM_181705 Homo sapiens hypothetical protein LOC90624 (LOC90624), mRNA
 NM_181706 Homo sapiens zinc finger, CSL domain containing 3 (ZC3L3), mRNA
 NM_181707 Homo sapiens hypothetical protein LOC124773 (LOC124773), mRNA
 NM_181708 Homo sapiens hypothetical protein LOC144233 (LOC144233), mRNA
 NM_181709 Homo sapiens hypothetical protein LOC144347 (LOC144347), mRNA
 NM_181710 Homo sapiens zinc and ring finger 4 (ZNRF4), mRNA
 NM_181711 Homo sapiens GRP1 (general receptor for phosphoinositides 1)-associated t
 NM_181712 Homo sapiens hypothetical protein LOC163782 (LOC163782), mRNA
 NM_181713 Homo sapiens UBXL domain containing 4 (UBXD4), mRNA
 NM_181714 Homo sapiens chromosome 6 open reading frame 152 (C6orf152), mRNA
 NM_181715 Homo sapiens transducer of regulated cAMP response element-binding prot
 NM_181716 Homo sapiens nuclear protein p30 (p30), mRNA
 NM_181717 Homo sapiens HLA complex group 27 (HCG27), mRNA
 NM_181718 Homo sapiens hypothetical protein LOC253982 (LOC253982), mRNA
 NM_181719 Homo sapiens hypothetical protein LOC255104 (LOC255104), mRNA

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NM_181720 Homo sapiens hypothetical protein LOC257106 (LOC257106), mRNA
 NM_181721 Homo sapiens forkhead box R1 (FOXR1), mRNA
 NM_181722 Homo sapiens hypothetical protein LOC285908 (LOC285908), mRNA
 NM_181723 Homo sapiens hypothetical protein LOC286097 (LOC286097), mRNA
 NM_181724 Homo sapiens hypothetical protein LOC338773 (LOC338773), mRNA
 NM_181725 Homo sapiens hypothetical protein FLJ12760 (FLJ12760), mRNA
 NM_181726 Homo sapiens low density lipoprotein receptor-related protein binding protein
 NM_181727 Homo sapiens spermatogenesis associated 12 (SPATA12), mRNA
 NM_181733 Homo sapiens component of oligomeric golgi complex 5 (COG5), transcript v
 NM_181737 Homo sapiens peroxiredoxin 2 (PRDX2), nuclear gene encoding mitochondri
 NM_181738 Homo sapiens peroxiredoxin 2 (PRDX2), nuclear gene encoding mitochondri
 NM_181739 Homo sapiens WINS1 protein with Drosophila Lines (Lin) homologous doma
 NM_181740 Homo sapiens WINS1 protein with Drosophila Lines (Lin) homologous doma
 NM_181741 Homo sapiens origin recognition complex, subunit 4-like (yeast) (ORC4L), tr
 NM_181742 Homo sapiens origin recognition complex, subunit 4-like (yeast) (ORC4L), tr
 NM_181744 Homo sapiens opsin 5 (OPN5), mRNA
 NM_181745 Homo sapiens G protein-coupled receptor 120 (GPR120), mRNA
 NM_181746 Homo sapiens LAG1 longevity assurance homolog 2 (S. cerevisiae) (LASS2)
 NM_181747 Homo sapiens origin recognition complex, subunit 5-like (yeast) (ORC5L), tr
 NM_181755 Homo sapiens hydroxysteroid (11-beta) dehydrogenase 1 (HSD11B1), transc
 NM_181756 Homo sapiens zinc finger protein 233 (ZNF233), mRNA
 NM_181762 Homo sapiens ubiquitin-conjugating enzyme E2A (RAD6 homolog) (UBE2A),
 NM_181773 Homo sapiens ARP10 protein (ARP10), mRNA
 NM_181774 Homo sapiens solute carrier family 36 (proton/amino acid symporter), memb
 NM_181775 Homo sapiens hypothetical protein DKFZp434G0625 (DKFZp434G0625), m
 NM_181776 Homo sapiens solute carrier family 36 (proton/amino acid symporter), memb
 NM_181777 Homo sapiens ubiquitin-conjugating enzyme E2A (RAD6 homolog) (UBE2A),
 NM_181780 Homo sapiens B and T lymphocyte associated (BTLA), mRNA
 NM_181781 Homo sapiens hypothetical protein FLJ20403 (FLJ20403), transcript variant :
 NM_181782 Homo sapiens nuclear receptor coactivator 7 (NCOA7), mRNA
 NM_181783 Homo sapiens SMILE protein (SMILE), mRNA
 NM_181784 Homo sapiens sprouty-related, EVH1 domain containing 2 (SPRED2), mRNA
 NM_181785 Homo sapiens hypothetical protein LOC283537 (LOC283537), mRNA
 NM_181786 Homo sapiens GLI-Kruppel family member HKR1 (HKR1), mRNA
 NM_181787 Homo sapiens hypothetical protein LOC286148 (LOC286148), mRNA
 NM_181788 Homo sapiens HANP1 (LOC341567), mRNA
 NM_181789 Homo sapiens collomin (COLM), mRNA
 NM_181790 Homo sapiens G protein-coupled receptor 142 (GPR142), mRNA
 NM_181791 Homo sapiens G protein-coupled receptor 141 (GPR141), mRNA
 NM_181794 Homo sapiens protein kinase (cAMP-dependent, catalytic) inhibitor beta (PKI
 NM_181795 Homo sapiens protein kinase (cAMP-dependent, catalytic) inhibitor beta (PKI
 NM_181797 Homo sapiens potassium voltage-gated channel, KQT-like subfamily, memb
 NM_181798 Homo sapiens potassium voltage-gated channel, KQT-like subfamily, memb
 NM_181799 Homo sapiens ubiquitin-conjugating enzyme E2C (UBE2C), transcript varian
 NM_181800 Homo sapiens ubiquitin-conjugating enzyme E2C (UBE2C), transcript varian
 NM_181801 Homo sapiens ubiquitin-conjugating enzyme E2C (UBE2C), transcript varian
 NM_181802 Homo sapiens ubiquitin-conjugating enzyme E2C (UBE2C), transcript varian
 NM_181803 Homo sapiens ubiquitin-conjugating enzyme E2C (UBE2C), transcript varian
 NM_181804 Homo sapiens protein kinase (cAMP-dependent, catalytic) inhibitor gamma (I
 NM_181805 Homo sapiens protein kinase (cAMP-dependent, catalytic) inhibitor gamma (I
 NM_181806 Homo sapiens 2-aminoadipic 6-semialdehyde dehydrogenase (NRPS998), n
 NM_181807 Homo sapiens doublecortin domain containing 1 (DCDC1), mRNA
 NM_181808 Homo sapiens polymerase (DNA directed) nu (POLN), mRNA
 NM_181809 Homo sapiens bone morphogenetic protein 8a (BMP8A), mRNA
 NM_181814 Homo sapiens chromosome 14 open reading frame 29 (C14orf29), transcript
 NM_181825 Homo sapiens neurofibromin 2 (bilateral acoustic neuroma) (NF2), transcript
 NM_181826 Homo sapiens neurofibromin 2 (bilateral acoustic neuroma) (NF2), transcript

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NM_181827 Homo sapiens neurofibromin 2 (bilateral acoustic neuroma) (NF2), transcript
 NM_181828 Homo sapiens neurofibromin 2 (bilateral acoustic neuroma) (NF2), transcript
 NM_181829 Homo sapiens neurofibromin 2 (bilateral acoustic neuroma) (NF2), transcript
 NM_181830 Homo sapiens neurofibromin 2 (bilateral acoustic neuroma) (NF2), transcript
 NM_181831 Homo sapiens neurofibromin 2 (bilateral acoustic neuroma) (NF2), transcript
 NM_181832 Homo sapiens neurofibromin 2 (bilateral acoustic neuroma) (NF2), transcript
 NM_181833 Homo sapiens neurofibromin 2 (bilateral acoustic neuroma) (NF2), transcript
 NM_181834 Homo sapiens neurofibromin 2 (bilateral acoustic neuroma) (NF2), transcript
 NM_181835 Homo sapiens neurofibromin 2 (bilateral acoustic neuroma) (NF2), transcript
 NM_181836 Homo sapiens CGI-109 protein (CGI-109), mRNA
 NM_181837 Homo sapiens origin recognition complex, subunit 3-like (yeast) (ORC3L), tr
 NM_181838 Homo sapiens ubiquitin-conjugating enzyme E2D 2 (UBC4/5 homolog, yeast
 NM_181839 Homo sapiens protein kinase (cAMP-dependent, catalytic) inhibitor alpha (PF
 NM_181840 Homo sapiens TWIK-related spinal cord K⁺ channel (TRIK), mRNA
 NM_181841 Homo sapiens transmembrane channel-like 3 (TMC3), mRNA
 NM_181842 Homo sapiens zinc finger and BTB domain containing 12 (ZBTB12), mRNA
 NM_181843 Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 8 (I
 NM_181844 Homo sapiens B-cell CLL/lymphoma 6, member B (zinc finger protein) (BCL6
 NM_181846 Homo sapiens GLI-Kruppel family member HKR2 (HKR2), mRNA
 NM_181847 Homo sapiens amphoterin induced gene 2 (AMIGO2), mRNA
 NM_181861 Homo sapiens apoptotic protease activating factor (APAF1), transcript varian
 NM_181862 Homo sapiens brain acyl-CoA hydrolase (BACH), transcript variant hBACHa/
 NM_181863 Homo sapiens brain acyl-CoA hydrolase (BACH), transcript variant hBACHa/
 NM_181864 Homo sapiens brain acyl-CoA hydrolase (BACH), transcript variant hBACHb,
 NM_181865 Homo sapiens brain acyl-CoA hydrolase (BACH), transcript variant hBACHc,
 NM_181866 Homo sapiens brain acyl-CoA hydrolase (BACH), transcript variant hBACHd,
 NM_181868 Homo sapiens apoptotic protease activating factor (APAF1), transcript varian
 NM_181869 Homo sapiens apoptotic protease activating factor (APAF1), transcript varian
 NM_181870 Homo sapiens dishevelled, dsh homolog 1 (Drosophila) (DVL1), transcript va
 NM_181871 Homo sapiens pyruvate kinase, liver and RBC (PKLR), nuclear gene encodir
 NM_181872 Homo sapiens doublesex and mab-3 related transcription factor 2 (DMRT2),
 NM_181873 Homo sapiens cisplatin resistance associated (CRA), mRNA
 NM_181874 Homo sapiens glutamate receptor, metabotropic 7 (GRM7), transcript variant
 NM_181875 Homo sapiens glutamate receptor, metabotropic 7 (GRM7), transcript variant
 NM_181876 Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit B (PF
 NM_181877 Homo sapiens zinc finger protein 29 (ZFP29), mRNA
 NM_181879 Homo sapiens leukocyte Ig-like receptor 9 (LIR9), transcript variant 3, mRNA
 NM_181880 Homo sapiens variable charge, Y-linked 1B (VCY1B), mRNA
 NM_181882 Homo sapiens perfaxin (PRX), mRNA
 NM_181885 Homo sapiens G protein-coupled receptor 100 (GPR100), mRNA
 NM_181886 Homo sapiens ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast
 NM_181887 Homo sapiens ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast
 NM_181888 Homo sapiens ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast
 NM_181889 Homo sapiens ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast
 NM_181890 Homo sapiens ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast
 NM_181891 Homo sapiens ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast
 NM_181892 Homo sapiens ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast
 NM_181893 Homo sapiens ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast
 NM_181894 Homo sapiens glutamate receptor, ionotropic, AMPA 3 (GRIA3), transcript v
 NM_181897 Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit B", al
 NM_181900 Homo sapiens START domain containing 5 (STARD5), transcript variant 1, n
 NM_181985 Homo sapiens leukocyte Ig-like receptor 9 (LIR9), transcript variant 2, mRNA
 NM_181986 Homo sapiens leukocyte Ig-like receptor 9 (LIR9), transcript variant 4, mRNA
 NM_182314 Homo sapiens cytosolic ovarian carcinoma antigen 1 (COVA1), transcript vai
 NM_182398 Homo sapiens ribosomal protein S6 kinase, 90kDa, polypeptide 5 (RPS6KAE
 NM_182470 Homo sapiens pyruvate kinase, muscle (PKM2), transcript variant 2, mRNA
 NM_182471 Homo sapiens pyruvate kinase, muscle (PKM2), transcript variant 3, mRNA

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NM_182472 Homo sapiens EphA5 (EPHA5), transcript variant 2, mRNA
 NM_182476 Homo sapiens coenzyme Q6 homolog (yeast) (COQ6), transcript variant 1, n
 NM_182477 Homo sapiens gametogenetin (GGN), transcript variant 2, mRNA
 NM_182480 Homo sapiens coenzyme Q6 homolog (yeast) (COQ6), transcript variant 2, n
 NM_182481 Homo sapiens B melanoma antigen family, member 3 (BAGE3), mRNA
 NM_182482 Homo sapiens B melanoma antigen family, member 2 (BAGE2), mRNA
 NM_182483 Homo sapiens NSFL1 (p97) cofactor (p47) (NSFL1C), transcript variant 3, m
 NM_182484 Homo sapiens B melanoma antigen family, member 5 (BAGE5), mRNA
 NM_182485 Homo sapiens cytoplasmic polyadenylation element binding protein 2 (CPEB
 NM_182486 Homo sapiens C1q and tumor necrosis factor related protein 6 (C1QTNF6), t
 NM_182487 Homo sapiens olfactomedin-like 2A (OLFM2A), mRNA
 NM_182488 Homo sapiens ubiquitin specific protease 12 (USP12), mRNA
 NM_182489 Homo sapiens STRA8 (LOC346873), mRNA
 NM_182490 Homo sapiens zinc finger protein 227 (ZNF227), mRNA
 NM_182491 Homo sapiens hypothetical protein LOC90637 (LOC90637), mRNA
 NM_182492 Homo sapiens hypothetical protein DKFZp434O0213 (DKFZp434O0213), mf
 NM_182493 Homo sapiens myosin light chain kinase (MLCK) (LOC91807), mRNA
 NM_182494 Homo sapiens family with sequence similarity 26, member A (FAM26A), mRf
 NM_182495 Homo sapiens hypothetical protein FLJ25224 (FLJ25224), mRNA
 NM_182496 Homo sapiens hypothetical protein FLJ40089 (FLJ40089), mRNA
 NM_182497 Homo sapiens type I hair keratin KA36 (KA36), mRNA
 NM_182498 Homo sapiens hypothetical protein MGC51082 (MGC51082), mRNA
 NM_182499 Homo sapiens hypothetical protein DKFZp434M202 (DKFZp434M202), mRN
 NM_182500 Homo sapiens hypothetical protein FLJ25143 (FLJ25143), mRNA
 NM_182501 Homo sapiens hypothetical protein MGC61716 (MGC61716), mRNA
 NM_182502 Homo sapiens hypothetical protein DKFZp686L1818 (DKFZp686L1818), mR
 NM_182503 Homo sapiens deaminase domain containing 1 (DEADC1), mRNA
 NM_182504 Homo sapiens Williams-Beuren syndrome critical region 28 (WBSCR28), mF
 NM_182505 Homo sapiens chromosome 9 open reading frame 85 (C9orf85), transcript v
 NM_182506 Homo sapiens hypothetical protein FLJ32965 (FLJ32965), mRNA
 NM_182507 Homo sapiens hypothetical protein LOC144501 (LOC144501), mRNA
 NM_182508 Homo sapiens hypothetical protein FLJ40919 (FLJ40919), mRNA
 NM_182509 Homo sapiens thrombospondin, type I, domain containing 3 (THSD3), transc
 NM_182510 Homo sapiens hypothetical protein FLJ32252 (FLJ32252), mRNA
 NM_182511 Homo sapiens cerebellin 2 precursor (CBLN2), mRNA
 NM_182513 Homo sapiens kinetochore protein Spc24 (Spc24), mRNA
 NM_182516 Homo sapiens hypothetical protein FLJ32011 (FLJ32011), mRNA
 NM_182517 Homo sapiens hypothetical protein MGC52423 (MGC52423), mRNA
 NM_182518 Homo sapiens hypothetical protein LOC149469 (LOC149469), mRNA
 NM_182519 Homo sapiens chromosome 20 open reading frame 186 (C20orf186), mRNA
 NM_182520 Homo sapiens chromosome 22 open reading frame 15 (C22orf15), mRNA
 NM_182521 Homo sapiens zinc finger, SWIM domain containing 2 (ZSWIM2), mRNA
 NM_182522 Homo sapiens TAF44 protein (TAF44), mRNA
 NM_182523 Homo sapiens hypothetical protein MGC61571 (MGC61571), mRNA
 NM_182524 Homo sapiens zinc finger protein 595 (ZNF595), mRNA
 NM_182525 Homo sapiens hypothetical protein FLJ32770 (FLJ32770), mRNA
 NM_182526 Homo sapiens hypothetical protein FLJ33387 (FLJ33387), mRNA
 NM_182527 Homo sapiens calcium binding protein 7 (CABP7), mRNA
 NM_182528 Homo sapiens complement component 1, q subcomponent-like 2 (C1QL2), r
 NM_182529 Homo sapiens THAP domain containing 5 (THAP5), mRNA
 NM_182530 Homo sapiens hypothetical protein FLJ25056 (FLJ25056), mRNA
 NM_182531 Homo sapiens hypothetical protein FLJ31875 (FLJ31875), mRNA
 NM_182532 Homo sapiens hypothetical protein LOC199964 (LOC199964), mRNA
 NM_182533 Homo sapiens hypothetical protein FLJ31031 (FLJ31031), mRNA
 NM_182534 Homo sapiens hypothetical protein FLJ23703 (FLJ23703), mRNA
 NM_182535 Homo sapiens hypothetical protein LOC200261 (LOC200261), mRNA
 NM_182536 Homo sapiens down-regulated in gastric cancer GDDR (GDDR), mRNA

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NM_182537 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 3 family member D (

NM_182538 Homo sapiens hypothetical protein MGC29671 (MGC29671), mRNA

NM_182539 Homo sapiens hypothetical protein MGC33600 (MGC33600), mRNA

NM_182541 Homo sapiens transmembrane protein 31 (TMEM31), mRNA

NM_182543 Homo sapiens nucleolar protein (NOL1/NOP2/sun) and PUA domains 1 (NO

NM_182546 Homo sapiens hypothetical protein MGC33530 (MGC33530), mRNA

NM_182547 Homo sapiens putative NFkB activating protein HNLF (HNLF), mRNA

NM_182548 Homo sapiens hypothetical protein MGC33835 (MGC33835), mRNA

NM_182549 Homo sapiens major histocompatibility complex, class II, DQ beta 2 (HLA-DQ

NM_182551 Homo sapiens acyl-CoA:lysocardiolipin acyltransferase 1 (ALCAT1), transcr

NM_182552 Homo sapiens hypothetical protein MGC43690 (MGC43690), mRNA

NM_182553 Homo sapiens hypothetical protein MGC50896 (MGC50896), mRNA

NM_182554 Homo sapiens chromosome 10 open reading frame 53 (C10orf53), mRNA

NM_182556 Homo sapiens hypothetical protein LOC283130 (LOC283130), mRNA

NM_182557 Homo sapiens B-cell CLL/lymphoma 9-like (BCL9L), mRNA

NM_182558 Homo sapiens hypothetical protein FLJ33810 (FLJ33810), mRNA

NM_182559 Homo sapiens hypothetical protein MGC57341 (MGC57341), mRNA

NM_182560 Homo sapiens hypothetical protein FLJ25773 (FLJ25773), mRNA

NM_182561 Homo sapiens hypothetical protein FLJ36144 (FLJ36144), mRNA

NM_182562 Homo sapiens hypothetical protein FLJ39743 (FLJ39743), mRNA

NM_182563 Homo sapiens hypothetical protein MGC21830 (MGC21830), mRNA

NM_182564 Homo sapiens hypothetical protein FLJ40319 (FLJ40319), mRNA

NM_182565 Homo sapiens hypothetical protein MGC29814 (MGC29814), mRNA

NM_182566 Homo sapiens secretory protein LOC284013 (LOC284013), mRNA

NM_182568 Homo sapiens hypothetical protein FLJ36492 (FLJ36492), mRNA

NM_182569 Homo sapiens hypothetical protein FLJ37451 (FLJ37451), mRNA

NM_182570 Homo sapiens hypothetical protein FLJ25715 (FLJ25715), mRNA

NM_182572 Homo sapiens zinc finger and SCAN domain containing 1 (ZSCAN1), mRNA

NM_182573 Homo sapiens hypothetical protein FLJ30469 (FLJ30469), mRNA

NM_182574 Homo sapiens hypothetical protein FLJ36070 (FLJ36070), mRNA

NM_182575 Homo sapiens hypothetical protein MGC34799 (MGC34799), mRNA

NM_182577 Homo sapiens chromosome 19 open reading frame 19 (C19orf19), mRNA

NM_182578 Homo sapiens hypothetical protein FLJ37964 (FLJ37964), mRNA

NM_182579 Homo sapiens hypothetical protein FLJ40343 (FLJ40343), mRNA

NM_182580 Homo sapiens cytochrome b-561 domain containing 1 (CYB561D1), mRNA

NM_182581 Homo sapiens hypothetical protein LOC284680 (LOC284680), mRNA

NM_182583 Homo sapiens hypothetical protein FLJ38374 (FLJ38374), mRNA

NM_182584 Homo sapiens hypothetical protein FLJ33706 (FLJ33706), mRNA

NM_182585 Homo sapiens hypothetical protein DKFZp451M2119 (DKFZp451M2119), m

NM_182586 Homo sapiens hypothetical protein FLJ33534 (FLJ33534), mRNA

NM_182587 Homo sapiens chromosome 2 open reading frame 21 (C2orf21), mRNA

NM_182589 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 3, family member E

NM_182590 Homo sapiens hypothetical protein FLJ33651 (FLJ33651), mRNA

NM_182591 Homo sapiens hypothetical protein FLJ37673 (FLJ37673), mRNA

NM_182592 Homo sapiens hypothetical protein FLJ39576 (FLJ39576), mRNA

NM_182594 Homo sapiens zinc finger protein 454 (ZNF454), mRNA

NM_182595 Homo sapiens hypothetical protein DKFZp564N2472 (DKFZp564N2472), m

NM_182596 Homo sapiens hypothetical protein FLJ25037 (FLJ25037), mRNA

NM_182597 Homo sapiens hypothetical protein FLJ39575 (FLJ39575), mRNA

NM_182598 Homo sapiens hypothetical protein FLJ36980 (FLJ36980), mRNA

NM_182600 Homo sapiens hypothetical protein LOC286359 (LOC286359), mRNA

NM_182603 Homo sapiens hypothetical protein FLJ37874 (FLJ37874), mRNA

NM_182605 Homo sapiens hypothetical protein FLJ40448 (FLJ40448), mRNA

NM_182606 Homo sapiens hypothetical protein LOC339967 (LOC339967), mRNA

NM_182607 Homo sapiens hypothetical protein MGC44287 (MGC44287), mRNA

NM_182608 Homo sapiens hypothetical protein DKFZp686O1689 (DKFZp686O1689), m

NM_182609 Homo sapiens hypothetical protein MGC48625 (MGC48625), mRNA

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NM_182610 Homo sapiens sterile alpha motif domain containing 7 (SAMD7), mRNA
 NM_182611 Homo sapiens G protein-coupled receptor 144 (GPR144), mRNA
 NM_182612 Homo sapiens hypothetical protein FLJ34283 (FLJ34283), mRNA
 NM_182613 Homo sapiens hypothetical protein FLJ33915 (FLJ33915), mRNA
 NM_182614 Homo sapiens hypothetical protein MGC20579 (MGC20579), mRNA
 NM_182615 Homo sapiens hypothetical protein MGC40069 (MGC40069), mRNA
 NM_182616 Homo sapiens hypothetical protein MGC61550 (MGC61550), mRNA
 NM_182617 Homo sapiens xenobiotic/medium-chain fatty acid:CoA ligase (HXMA), nucle
 NM_182619 Homo sapiens secretory protein LOC348174 (LOC348174), mRNA
 NM_182620 Homo sapiens family with sequence similarity 33, member A (FAM33A), mR
 NM_182621 Homo sapiens hypothetical protein MGC52498 (MGC52498), mRNA
 NM_182623 Homo sapiens hypothetical protein FLJ36766 (FLJ36766), mRNA
 NM_182625 Homo sapiens hypothetical protein FLJ40869 (FLJ40869), mRNA
 NM_182626 Homo sapiens hypothetical protein FLJ25102 (FLJ25102), mRNA
 NM_182627 Homo sapiens hypothetical protein MGC64882 (MGC64882), mRNA
 NM_182628 Homo sapiens hypothetical protein FLJ40083 (FLJ40083), mRNA
 NM_182631 Homo sapiens hypothetical protein LOC348840 (LOC348840), mRNA
 NM_182632 Homo sapiens solute carrier family 6 (neurotransmitter transporter), member
 NM_182633 Homo sapiens hypothetical protein FLJ39963 (FLJ39963), mRNA
 NM_182634 Homo sapiens hypothetical protein FLJ36166 (FLJ36166), mRNA
 NM_182635 Homo sapiens hypothetical protein LOC349236 (LOC349236), mRNA
 NM_182637 Homo sapiens Hermansky-Pudlak syndrome 1 (HPS1), transcript variant 2, r
 NM_182638 Homo sapiens Hermansky-Pudlak syndrome 1 (HPS1), transcript variant 4, r
 NM_182639 Homo sapiens Hermansky-Pudlak syndrome 1 (HPS1), transcript variant 3, r
 NM_182640 Homo sapiens mitochondrial ribosomal protein S9 (MRPS9), nuclear gene er
 NM_182641 Homo sapiens fetal Alzheimer antigen (FALZ), transcript variant 1, mRNA
 NM_182642 Homo sapiens CTD (carboxy-terminal domain, RNA polymerase II, polypepti
 NM_182643 Homo sapiens deleted in liver cancer 1 (DLC1), transcript variant 1, mRNA
 NM_182644 Homo sapiens EphA3 (EPHA3), transcript variant 2, mRNA
 NM_182645 Homo sapiens vestigial like 2 (Drosophila) (VGLL2), transcript variant 1, mR
 NM_182646 Homo sapiens cytoplasmic polyadenylation element binding protein 2 (CPEB
 NM_182647 Homo sapiens opiate receptor-like 1 (OPRL1), transcript variant 1, mRNA
 NM_182648 Homo sapiens bromodomain adjacent to zinc finger domain, 1A (BAZ1A), tra
 NM_182649 Homo sapiens proliferating cell nuclear antigen (PCNA), transcript variant 2,
 NM_182658 Homo sapiens chromosome 20 open reading frame 185 (C20orf185), mRNA
 NM_182659 Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, Y-link
 NM_182660 Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, Y-link
 NM_182661 Homo sapiens ceramide kinase (CERK), transcript variant 2, mRNA
 NM_182662 Homo sapiens aminoadipate aminotransferase (AADAT), transcript variant 2.
 NM_182663 Homo sapiens Ras association (RalGDS/AF-6) domain family 5 (RASSF5), t
 NM_182664 Homo sapiens Ras association (RalGDS/AF-6) domain family 5 (RASSF5), t
 NM_182665 Homo sapiens Ras association (RalGDS/AF-6) domain family 5 (RASSF5), t
 NM_182666 Homo sapiens ubiquitin-conjugating enzyme E2E 1 (UBC4/5 homolog, yeast
 NM_182676 Homo sapiens phospholipid transfer protein (PLTP), transcript variant 2, mR
 NM_182678 Homo sapiens ubiquitin-conjugating enzyme E2E 3 (UBC4/5 homolog, yeast
 NM_182679 Homo sapiens hypothetical protein FLJ20249 (FLJ20249), transcript variant
 NM_182680 Homo sapiens amelogenin (amelogenesis imperfecta 1, X-linked) (AMELX),
 NM_182681 Homo sapiens amelogenin (amelogenesis imperfecta 1, X-linked) (AMELX),
 NM_182682 Homo sapiens ubiquitin-conjugating enzyme E2G 1 (UBC7 homolog, C. eleg
 NM_182683 Homo sapiens uroplakin 3B (UPK3B), transcript variant 3, mRNA
 NM_182684 Homo sapiens uroplakin 3B (UPK3B), transcript variant 2, mRNA
 NM_182685 Homo sapiens ephrin-A1 (EFNA1), transcript variant 2, mRNA
 NM_182686 Homo sapiens polycystic kidney disease 1-like (PKD1-like), transcript variant
 NM_182687 Homo sapiens membrane-associated tyrosine- and threonine-specific cdc2-li
 NM_182688 Homo sapiens ubiquitin-conjugating enzyme E2G 2 (UBC7 homolog, yeast) (i
 NM_182689 Homo sapiens ephrin-A4 (EFNA4), transcript variant 2, mRNA
 NM_182690 Homo sapiens ephrin-A4 (EFNA4), transcript variant 3, mRNA

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NM_182691 Homo sapiens SFRS protein kinase 2 (SRPK2), transcript variant 2, mRNA
 NM_182692 Homo sapiens SFRS protein kinase 2 (SRPK2), transcript variant 1, mRNA
 NM_182697 Homo sapiens ubiquitin-conjugating enzyme E2H (UBC8 homolog, yeast) (U
 NM_182699 Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 53 (DXX53), mRNA
 NM_182700 Homo sapiens Sp8 transcription factor (SP8), transcript variant 1, mRNA
 NM_182701 Homo sapiens glutathione peroxidase 6 (olfactory) (GPX6), mRNA
 NM_182702 Homo sapiens testis serine protease 2 (TESSP2), mRNA
 NM_182703 Homo sapiens hypothetical protein LOC348094 (LOC348094), mRNA
 NM_182704 Homo sapiens selenoprotein V (SELV), mRNA
 NM_182705 Homo sapiens hypothetical protein MGC45871 (MGC45871), mRNA
 NM_182706 Homo sapiens scribbled homolog (Drosophila) (SCRIB), transcript variant 1,
 NM_182709 Homo sapiens HIV-1 Tat interacting protein, 60kDa (HTATIP), transcript vari
 NM_182710 Homo sapiens HIV-1 Tat interacting protein, 60kDa (HTATIP), transcript vari
 NM_182712 Homo sapiens eukaryotic translation initiation factor 3, subunit 9 eta, 116kDa
 NM_182715 Homo sapiens synaptophysin-like protein (SYPL), transcript variant 2, mRNA
 NM_182717 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_182718 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_182719 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_182720 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_182721 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_182722 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_182723 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_182724 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_182725 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_182728 Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+ sys
 NM_182729 Homo sapiens thioredoxin reductase 1 (TXNRD1), transcript variant 5, mRNA
 NM_182734 Homo sapiens phospholipase C, beta 1 (phosphoinositide-specific) (PLCB1),
 NM_182739 Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 6, 17
 NM_182740 Homo sapiens polycystic kidney disease 1-like 2 (PKD1L2), transcript varian
 NM_182741 Homo sapiens mucin 1, transmembrane (MUC1), mRNA
 NM_182742 Homo sapiens thioredoxin reductase 1 (TXNRD1), transcript variant 2, mRNA
 NM_182743 Homo sapiens thioredoxin reductase 1 (TXNRD1), transcript variant 4, mRNA
 NM_182744 Homo sapiens neuroblastoma, suppression of tumorigenicity 1 (NBL1), trans
 NM_182746 Homo sapiens MCM4 minichromosome maintenance deficient 4 (S. cerevisi
 NM_182749 Homo sapiens chromosome 21 open reading frame 127 (C21orf127), transcr
 NM_182751 Homo sapiens MCM10 minichromosome maintenance deficient 10 (S. cerevi
 NM_182752 Homo sapiens hypothetical protein LOC127262 (LOC127262), mRNA
 NM_182755 Homo sapiens hypothetical protein LOC220929 (LOC220929), mRNA
 NM_182756 Homo sapiens speedy homolog 1 (Drosophila) (SPDY1), mRNA
 NM_182757 Homo sapiens IBR domain containing 2 (IBRDC2), mRNA
 NM_182758 Homo sapiens hypothetical protein FLJ38736 (FLJ38736), mRNA
 NM_182759 Homo sapiens TAF3 protein (TAF3), mRNA
 NM_182760 Homo sapiens sulfatase modifying factor 1 (SUMF1), mRNA
 NM_182761 Homo sapiens hypothetical protein LOC340069 (LOC340069), mRNA
 NM_182762 Homo sapiens putative binding protein 7a5 (7A5), mRNA
 NM_182763 Homo sapiens myeloid cell leukemia sequence 1 (BCL2-related) (MCL1), tra
 NM_182764 Homo sapiens engulfment and cell motility 2 (ced-12 homolog, C. elegans) (I
 NM_182765 Homo sapiens HECT domain containing 2 (HECTD2), transcript variant 1, ml
 NM_182766 Homo sapiens hypothetical protein FLJ32940 (DKFZp686H1423), transcript
 NM_182767 Homo sapiens solute carrier family 6 (neurotransmitter transporter), member
 NM_182769 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_182770 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_182771 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_182772 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_182774 Homo sapiens hypothetical protein LOC259173 (FLJ36525), transcript varian
 NM_182775 Homo sapiens hypothetical protein LOC259173 (FLJ36525), transcript varian
 NM_182776 Homo sapiens MCM7 minichromosome maintenance deficient 7 (S. cerevisi

NM_182777 Homo sapiens ribosomal protein S3A (RPS3A), mRNA
 NM_182779 Homo sapiens dishevelled, dsh homolog 1 (Drosophila) (DVL1), mRNA
 NM_182789 Homo sapiens poly(A) binding protein interacting protein 1 (PAIP1), transcript
 NM_182790 Homo sapiens pre-B-cell colony enhancing factor 1 (PBEF1), transcript varia
 NM_182791 Homo sapiens hypothetical protein FLJ32855 (FLJ32855), mRNA
 NM_182792 Homo sapiens thymosin-like 1 (TMSL1), mRNA
 NM_182793 Homo sapiens thymosin-like 2 (TMSL2), mRNA
 NM_182794 Homo sapiens thymosin-like 4 (TMSL4), mRNA
 NM_182795 Homo sapiens nucleophosmin/nucleoplasmin, 2 (NPM2), mRNA
 NM_182796 Homo sapiens methionine adenosyltransferase II, beta (MAT2B), transcript v
 NM_182797 Homo sapiens phospholipase C, beta 4 (PLCB4), transcript variant 2, mRNA
 NM_182798 Homo sapiens hypothetical protein FLJ39155 (FLJ39155), transcript variant.
 NM_182799 Homo sapiens hypothetical protein FLJ39155 (FLJ39155), transcript variant.
 NM_182800 Homo sapiens arsenate resistance protein ARSZ (ARS2), transcript variant 2
 NM_182801 Homo sapiens hypothetical protein FLJ39155 (FLJ39155), transcript variant.
 NM_182802 Homo sapiens MCM8 minichromosome maintenance deficient 8 (S. cerevisia
 NM_182804 Homo sapiens apolipoprotein B48 receptor (APOB48R), mRNA
 NM_182810 Homo sapiens activating transcription factor 4 (tax-responsive enhancer elen
 NM_182811 Homo sapiens phospholipase C, gamma 1 (PLCG1), transcript variant 2, mR
 NM_182812 Homo sapiens splicing factor 4 (SF4), transcript variant c, mRNA
 NM_182826 Homo sapiens scavenger receptor class A, member 3 (SCARA3), transcript v
 NM_182827 Homo sapiens FK506 binding protein 9-like (FKBP9L), mRNA
 NM_182828 Homo sapiens growth differentiation factor 7 (GDF7), mRNA
 NM_182829 Homo sapiens hypothetical protein LOC158160 (LOC158160), mRNA
 NM_182830 Homo sapiens MAM domain containing 1 (MAMDC1), mRNA
 NM_182831 Homo sapiens TNT protein (TNT), mRNA
 NM_182832 Homo sapiens placenta-specific 4 (PLAC4), mRNA
 NM_182833 Homo sapiens GDP domain containing protein (LOC220032), mRNA
 NM_182835 Homo sapiens sec1 family domain containing 1 (SCFD1), transcript variant 2
 NM_182836 Homo sapiens Rab geranylgeranyltransferase, alpha subunit (RABGGTA), tr
 NM_182838 Homo sapiens solute carrier family 35, member E2 (SLC35E2), mRNA
 NM_182847 Homo sapiens amiloride-sensitive cation channel 4, pituitary (ACCN4), trans
 NM_182848 Homo sapiens claudin 10 (CLDN10), transcript variant 1, mRNA
 NM_182849 Homo sapiens cyclin B1 interacting protein 1 (CCNB1IP1), transcript variant.
 NM_182850 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_182851 Homo sapiens cyclin B1 interacting protein 1 (CCNB1IP1), transcript variant.
 NM_182852 Homo sapiens cyclin B1 interacting protein 1 (CCNB1IP1), transcript variant.
 NM_182853 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_182854 Homo sapiens selectin ligand interactor cytoplasmic-1 (SLIC1), mRNA
 NM_182894 Homo sapiens ceh-10 homeo domain containing homolog (C. elegans) (CHX
 NM_182895 Homo sapiens scavenger receptor class F, member 2 (SCARF2), transcript v
 NM_182896 Homo sapiens hypothetical protein DKFzp761H079 (DKFzp761H079), trans
 NM_182898 Homo sapiens cAMP responsive element binding protein 5 (CREB5), mRNA
 NM_182899 Homo sapiens cAMP responsive element binding protein 5 (CREB5), mRNA
 NM_182901 Homo sapiens chromosome 11 open reading frame 17 (C11orf17), transcript
 NM_182902 Homo sapiens kinesin family member 9 (KIF9), mRNA
 NM_182903 Homo sapiens kinesin family member 9 (KIF9), mRNA
 NM_182904 Homo sapiens procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-
 NM_182905 Homo sapiens CXYorf1-related protein (DKFzp434K1323), mRNA
 NM_182906 Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain
 NM_182907 Homo sapiens PR domain containing 1, with ZNF domain (PRDM1), transcript
 NM_182908 Homo sapiens dehydrogenase/reductase (SDR family) member 2 (DHRS2).
 NM_182909 Homo sapiens downregulated in ovarian cancer 1 (DOC1), transcript variant
 NM_182910 Homo sapiens spectrin repeat containing, nuclear envelope 2 (SYNE2), trans
 NM_182911 Homo sapiens testis specific, 10 (TSGA10), mRNA
 NM_182912 Homo sapiens spectrin repeat containing, nuclear envelope 2 (SYNE2), trans
 NM_182913 Homo sapiens spectrin repeat containing, nuclear envelope 2 (SYNE2), trans

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NM_182914 Homo sapiens spectrin repeat containing, nuclear envelope 2 (SYNE2), trans
 NM_182915 Homo sapiens dudulin 2 (TSAP6), mRNA
 NM_182916 Homo sapiens tRNA nucleotidyl transferase, CCA-adding, 1 (TRNT1), mRNA
 NM_182917 Homo sapiens eukaryotic translation initiation factor 4 gamma, 1 (EIF4G1), t
 NM_182918 Homo sapiens v-ets erythroblastosis virus E26 oncogene like (avian) (ERG),
 NM_182919 Homo sapiens TIR domain containing adaptor inducing interferon-beta (TRIF)
 NM_182920 Homo sapiens a disintegrin-like and metalloprotease (repolysin type) with th
 NM_182921 Homo sapiens a disintegrin-like and metalloprotease (repolysin type) with th
 NM_182922 Homo sapiens hypothetical protein FLJ20718 (FLJ20718), mRNA
 NM_182923 Homo sapiens kinesin 2 60/70kDa (KNS2), mRNA
 NM_182924 Homo sapiens MICAL-like 2 (FLJ23471), transcript variant 1, mRNA
 NM_182925 Homo sapiens fms-related tyrosine kinase 4 (FLT4), transcript variant 1, mRI
 NM_182926 Homo sapiens kinectin 1 (kinesin receptor) (KTN1), mRNA
 NM_182931 Homo sapiens myeloid/lymphoid or mixed-lineage leukemia 5 (trithorax hom
 NM_182932 Homo sapiens solute carrier family 8 (sodium-calcium exchanger), member 5
 NM_182933 Homo sapiens solute carrier family 8 (sodium-calcium exchanger), member 5
 NM_182934 Homo sapiens myelin-associated oligodendrocyte basic protein (MOBP), mR
 NM_182935 Homo sapiens myelin-associated oligodendrocyte basic protein (MOBP), mR
 NM_182936 Homo sapiens solute carrier family 8 (sodium-calcium exchanger), member 5
 NM_182943 Homo sapiens procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine hydr
 NM_182944 Homo sapiens ninein (GSK3B interacting protein) (NIN), transcript variant 1,
 NM_182945 Homo sapiens ninein (GSK3B interacting protein) (NIN), transcript variant 3,
 NM_182946 Homo sapiens ninein (GSK3B interacting protein) (NIN), transcript variant 5,
 NM_182947 Homo sapiens RAC/CDC42 exchange factor (GEFT), transcript variant 1, mR
 NM_182948 Homo sapiens protein kinase, cAMP-dependent, catalytic, beta (PRKACB), t
 NM_182960 Homo sapiens hypothetical protein MGC21644 (MGC21644), transcript varia
 NM_182961 Homo sapiens spectrin repeat containing, nuclear envelope 1 (SYNE1), trans
 NM_182962 Homo sapiens baculoviral IAP repeat-containing 3 (BIRC3), transcript variant
 NM_182964 Homo sapiens neuron navigator 2 (NAV2), transcript variant 1, mRNA
 NM_182965 Homo sapiens sphingosine kinase 1 (SPHK1), mRNA
 NM_182966 Homo sapiens neural precursor cell expressed, developmentally down-regul
 NM_182970 Homo sapiens regulating synaptic membrane exocytosis 4 (RIMS4), mRNA
 NM_182971 Homo sapiens cytochrome c oxidase subunit 8C (COX8C), mRNA
 NM_182972 Homo sapiens interferon regulatory factor 2 binding protein 2 (IRF2BP2), mR
 NM_182973 Homo sapiens transmembrane serine protease 9 (TMPRSS9), mRNA
 NM_182974 Homo sapiens galactosyltransferase family 6 domain containing 1 (GLTDC1)
 NM_182975 Homo sapiens hypothetical protein FLJ20403 (FLJ20403), transcript variant :
 NM_182976 Homo sapiens hypothetical protein FLJ20403 (FLJ20403), transcript variant :
 NM_182977 Homo sapiens nicotinamide nucleotide transhydrogenase (NNT), mRNA
 NM_182978 Homo sapiens guanine nucleotide binding protein (G protein), alpha activatin
 NM_182980 Homo sapiens pregnancy-induced growth inhibitor (OKL38), mRNA
 NM_182981 Homo sapiens pregnancy-induced growth inhibitor (OKL38), mRNA
 NM_182982 Homo sapiens G protein-coupled receptor kinase 4 (GRK4), transcript varian
 NM_182983 Homo sapiens hepsin (transmembrane protease, serine 1) (HPN), transcript
 NM_182984 Homo sapiens HpaII tiny fragments locus 9C (HTF9C), transcript variant 1, r
 NM_182985 Homo sapiens ring finger protein 36 (RNF36), transcript variant a, mRNA
 NM_183001 Homo sapiens SHC (Src homology 2 domain containing) transforming protei
 NM_183002 Homo sapiens solute carrier family 8 (sodium-calcium exchanger), member 5
 NM_183003 Homo sapiens cytochrome c oxidase subunit VIIa polypeptide 3 (liver) (COX)
 NM_183004 Homo sapiens eukaryotic translation initiation factor 5 (EIF5), transcript varia
 NM_183005 Homo sapiens ribonuclease P/MRP 38kDa subunit (RPP38), transcript varia
 NM_183006 Homo sapiens discs, large (Drosophila) homolog-associated protein 4 (DLG4)
 NM_183008 Homo sapiens socius (SOC), mRNA
 NM_183009 Homo sapiens DKFZP434116 protein (DKFZP434116), transcript variant 2,
 NM_183010 Homo sapiens trinucleotide repeat containing 5 (TNRCS), mRNA
 NM_183011 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_183012 Homo sapiens cAMP responsive element modulator (CREM), transcript varia

NM_183013 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_183040 Homo sapiens dystrobrevin binding protein 1 (DTNBP1), transcript variant 2,
 NM_183041 Homo sapiens dystrobrevin binding protein 1 (DTNBP1), transcript variant 3,
 NM_183043 Homo sapiens ring finger protein (C3H2C3 type) 6 (RNF6), transcript variant
 NM_183044 Homo sapiens ring finger protein (C3H2C3 type) 6 (RNF6), transcript variant
 NM_183045 Homo sapiens ring finger protein (C3H2C3 type) 6 (RNF6), transcript variant
 NM_183047 Homo sapiens protein kinase C binding protein 1 (PRKCBP1), transcript vari
 NM_183048 Homo sapiens protein kinase C binding protein 1 (PRKCBP1), transcript vari
 NM_183049 Homo sapiens thymosin-like 3 (TM3L3), mRNA
 NM_183050 Homo sapiens branched chain keto acid dehydrogenase E1, beta polypeptid
 NM_183057 Homo sapiens vacuolar protein sorting 28 (yeast) (VPS28), transcript variant
 NM_183058 Homo sapiens lysozyme-like 2 (LYL2), mRNA
 NM_183059 Homo sapiens chromosome 1 open reading frame 36 (C1orf36), mRNA
 NM_183060 Homo sapiens cAMP responsive element modulator (CREM), transcript varia
 NM_183062 Homo sapiens marapsin 2 (MPN2), mRNA
 NM_183063 Homo sapiens ring finger protein 7 (RNF7), transcript variant 2, mRNA
 NM_183065 Homo sapiens hypothetical protein MGC10744 (MGC10744), transcript varia
 NM_183075 Homo sapiens cytochrome P450, family 2, subfamily U, polypeptide 1 (CYP2
 NM_183078 Homo sapiens ring finger protein (C3HC4 type) 8 (RNF8), transcript variant 2
 NM_183079 Homo sapiens prion protein (p27-30) (Creutzfeld-Jakob disease, Gerstmann-
 NM_183227 Homo sapiens RAB23, member RAS oncogene family (RAB23), transcript ve
 NM_183228 Homo sapiens zinc finger protein, subfamily 1A, 3 (Aiolos) (ZNFN1A3), trans
 NM_183229 Homo sapiens zinc finger protein, subfamily 1A, 3 (Aiolos) (ZNFN1A3), trans
 NM_183230 Homo sapiens zinc finger protein, subfamily 1A, 3 (Aiolos) (ZNFN1A3), trans
 NM_183231 Homo sapiens zinc finger protein, subfamily 1A, 3 (Aiolos) (ZNFN1A3), trans
 NM_183232 Homo sapiens zinc finger protein, subfamily 1A, 3 (Aiolos) (ZNFN1A3), trans
 NM_183233 Homo sapiens solute carrier family 22 (organic cation transporter), member 1
 NM_183234 Homo sapiens RAB27A, member RAS oncogene family (RAB27A), transcript
 NM_183235 Homo sapiens RAB27A, member RAS oncogene family (RAB27A), transcript
 NM_183236 Homo sapiens RAB27A, member RAS oncogene family (RAB27A), transcript
 NM_183237 Homo sapiens ring finger protein 7 (RNF7), transcript variant 3, mRNA
 NM_183238 Homo sapiens zinc finger protein 605 (ZNF605), mRNA
 NM_183239 Homo sapiens glutathione S-transferase omega 2 (GSTO2), mRNA
 NM_183240 Homo sapiens voltage-dependent calcium channel gamma subunit-like prote
 NM_183241 Homo sapiens hypothetical protein LOC286257 (LOC286257), mRNA
 NM_183242 Homo sapiens BTB (POZ) domain containing 8 (BTBD8), mRNA
 NM_183243 Homo sapiens IMP (inosine monophosphate) dehydrogenase 1 (IMPDH1), tr
 NM_183244 Homo sapiens phosphatase and actin regulator 3 (PHACTR3), transcript vari
 NM_183245 Homo sapiens inversin (INVS), transcript variant 2, mRNA
 NM_183246 Homo sapiens phosphatase and actin regulator 3 (PHACTR3), transcript vari
 NM_183247 Homo sapiens transmembrane protease, serine 4 (TMPRSS4), transcript var
 NM_183323 Homo sapiens poly(A) binding protein interacting protein 1 (PAIP1), transcrip
 NM_183337 Homo sapiens regulator of G-protein signalling 11 (RGS11), transcript varian
 NM_183352 Homo sapiens SEC13-like 1 (S. cerevisiae) (SEC13L1), transcript variant 2, i
 NM_183353 Homo sapiens ring finger protein 12 (RNF12), transcript variant 2, mRNA
 NM_183356 Homo sapiens asparagine synthetase (ASNS), transcript variant 3, mRNA
 NM_183357 Homo sapiens adenylate cyclase 5 (ADCY5), mRNA
 NM_183359 Homo sapiens bromodomain containing 8 (BRD8), transcript variant 3, mRN
 NM_183360 Homo sapiens dystrobrevin, beta (DTNB), transcript variant 4, mRNA
 NM_183361 Homo sapiens dystrobrevin, beta (DTNB), transcript variant 5, mRNA
 NM_183372 Homo sapiens hypothetical protein LOC200030 (LOC200030), mRNA
 NM_183373 Homo sapiens chromosome 6 open reading frame 145 (C6orf145), mRNA
 NM_183374 Homo sapiens cytochrome P450, family 26, subfamily C, polypeptide 1 (CYP
 NM_183375 Homo sapiens epidermis-specific serine protease-like protein (ESSPL), mRN
 NM_183376 Homo sapiens arrestin domain containing 4 (ARRDC4), mRNA
 NM_183377 Homo sapiens amiloride-sensitive cation channel 1, neuronal (degenerin) (At
 NM_183378 Homo sapiens ovochymase 1 (OVCH1), mRNA

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NM_183379 Homo sapiens testis serine protease 1 (TESSP1), mRNA
 NM_183380 Homo sapiens dystonin (DST), transcript variant 1, mRNA
 NM_183381 Homo sapiens ring finger protein 13 (RNF13), transcript variant 4, mRNA
 NM_183382 Homo sapiens ring finger protein 13 (RNF13), transcript variant 2, mRNA
 NM_183383 Homo sapiens ring finger protein 13 (RNF13), transcript variant 3, mRNA
 NM_183384 Homo sapiens ring finger protein 13 (RNF13), transcript variant 5, mRNA
 NM_183385 Homo sapiens peroxisomal acyl-CoA thioesterase (PTE1), transcript variant:
 NM_183386 Homo sapiens peroxisomal acyl-CoA thioesterase (PTE1), transcript variant:
 NM_183387 Homo sapiens echinoderm microtubule associated protein like 5 (EML5), mR
 NM_183393 Homo sapiens Ca2+-dependent secretion activator (CADPS), transcript varie
 NM_183394 Homo sapiens Ca2+-dependent secretion activator (CADPS), transcript varie
 NM_183395 Homo sapiens cold autoinflammatory syndrome 1 (CIAS1), transcript variant
 NM_183397 Homo sapiens peroxisomal membrane protein 4, 24kDa (PXMP4), transcript
 NM_183398 Homo sapiens ring finger protein 14 (RNF14), transcript variant 2, mRNA
 NM_183399 Homo sapiens ring finger protein 14 (RNF14), transcript variant 3, mRNA
 NM_183400 Homo sapiens ring finger protein 14 (RNF14), transcript variant 4, mRNA
 NM_183401 Homo sapiens ring finger protein 14 (RNF14), transcript variant 5, mRNA
 NM_183404 Homo sapiens retinoblastoma-like 1 (p107) (RBL1), transcript variant 2, mR
 NM_183412 Homo sapiens F-box protein 44 (FBXO44), transcript variant 2, mRNA
 NM_183413 Homo sapiens F-box protein 44 (FBXO44), transcript variant 3, mRNA
 NM_183414 Homo sapiens ubiquitin protein ligase E3B (UBE3B), transcript variant 2, mR
 NM_183415 Homo sapiens ubiquitin protein ligase E3B (UBE3B), transcript variant 3, mR
 NM_183416 Homo sapiens kinesin family member 1B (KIF1B), transcript variant 2, mRN
 NM_183418 Homo sapiens molybdenum cofactor synthesis 2 (MOCS2), transcript variant
 NM_183419 Homo sapiens ring finger protein 19 (RNF19), transcript variant 1, mRNA
 NM_183420 Homo sapiens F-box protein 25 (FBXO25), transcript variant 2, mRNA
 NM_183421 Homo sapiens F-box protein 25 (FBXO25), transcript variant 1, mRNA
 NM_183422 Homo sapiens transforming growth factor beta 1 induced transcript 4 (TGFB
 NM_183425 Homo sapiens RNA-binding region (RNP1, RRM) containing 1 (RNPC1), trar
 NM_184041 Homo sapiens aldolase A, fructose-bisphosphate (ALDOA), transcript variant
 NM_184042 Homo sapiens Cohen syndrome 1 (COH1), transcript variant 2, mRNA
 NM_184043 Homo sapiens aldolase A, fructose-bisphosphate (ALDOA), transcript variant
 NM_184085 Homo sapiens ring finger protein 29 (RNF29), transcript variant 1, mRNA
 NM_184086 Homo sapiens ring finger protein 29 (RNF29), transcript variant 3, mRNA
 NM_184087 Homo sapiens ring finger protein 29 (RNF29), transcript variant 4, mRNA
 NM_184231 Homo sapiens NCK interacting protein with SH3 domain (NCKIPSD), transcr
 NM_184234 Homo sapiens RNA-binding region (RNP1, RRM) containing 2 (RNPC2), trar
 NM_184237 Homo sapiens RNA-binding region (RNP1, RRM) containing 2 (RNPC2), trar
 NM_184241 Homo sapiens RNA-binding region (RNP1, RRM) containing 2 (RNPC2), trar
 NM_184244 Homo sapiens RNA-binding region (RNP1, RRM) containing 2 (RNPC2), trar
 NM_187841 Homo sapiens ring finger protein 30 (RNF30), transcript variant 2, mRNA
 NM_194071 Homo sapiens cAMP responsive element binding protein 3-like 2 (CREB3L2)
 NM_194072 Homo sapiens spermatid-specific linker histone H1-like protein (HILS1), mR
 NM_194247 Homo sapiens heterogeneous nuclear ribonucleoprotein A3 (HNRPA3), mR
 NM_194248 Homo sapiens otoferlin (OTOF), transcript variant 1, mRNA
 NM_194249 Homo sapiens dead end homolog 1 (zebrafish) (DND1), mRNA
 NM_194250 Homo sapiens similar to C630007C17Rik protein (LOC91752), mRNA
 NM_194251 Homo sapiens G protein-coupled receptor 151 (GPR151), mRNA
 NM_194252 Homo sapiens chromosome 9 open reading frame 20 (C9orf20), mRNA
 NM_194255 Homo sapiens solute carrier family 19 (folate transporter), member 1 (SLC19
 NM_194259 Homo sapiens ubiquitin-conjugating enzyme E21 (UBC9 homolog, yeast) (UE
 NM_194260 Homo sapiens ubiquitin-conjugating enzyme E21 (UBC9 homolog, yeast) (UE
 NM_194261 Homo sapiens ubiquitin-conjugating enzyme E21 (UBC9 homolog, yeast) (UE
 NM_194270 Homo sapiens protein containing single MORN motif in testis (MOPT), mRN/
 NM_194271 Homo sapiens ring finger protein 34 (RNF34), transcript variant 1, mRNA
 NM_194276 Homo sapiens hypothetical protein FLJ20209 (FLJ20209), mRNA
 NM_194277 Homo sapiens hypothetical protein LOC90167 (LOC90167), mRNA

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NM_194278 Homo sapiens chromosome 14 open reading frame 43 (C14orf43), mRNA
 NM_194279 Homo sapiens HESB like domain containing 1 (HBLD1), mRNA
 NM_194281 Homo sapiens chromosome 18 open reading frame 37 (C18orf37), mRNA
 NM_194282 Homo sapiens hypothetical protein DKFZp686L1814 (DKFZp686L1814), mR
 NM_194283 Homo sapiens hypothetical protein LOC134218 (LOC134218), mRNA
 NM_194284 Homo sapiens claudin 23 (CLDN23), mRNA
 NM_194285 Homo sapiens hypothetical protein FLJ39441 (FLJ39441), mRNA
 NM_194286 Homo sapiens KIAA1853 protein (KIAA1853), mRNA
 NM_194287 Homo sapiens chromosome 14 open reading frame 166B (C14orf166B), mR
 NM_194288 Homo sapiens hypothetical protein LOC146712 (LOC146712), mRNA
 NM_194289 Homo sapiens hypothetical protein LOC152195 (LOC152195), mRNA
 NM_194290 Homo sapiens hypothetical protein LOC153684 (LOC153684), mRNA
 NM_194291 Homo sapiens hypothetical protein BC017881 (LOC157378), mRNA
 NM_194292 Homo sapiens hypothetical protein DKFZp761A078 (DKFZp761A078), mRN
 NM_194293 Homo sapiens cardiomyopathy associated 1 (CMYA1), mRNA
 NM_194294 Homo sapiens hypothetical protein LOC169355 (LOC169355), mRNA
 NM_194295 Homo sapiens hypothetical protein DKFZp434I020 (DKFZp434I020), mR
 NM_194298 Homo sapiens solute carrier family 16 (monocarboxylic acid transporters), m
 NM_194299 Homo sapiens hypothetical protein LOC221711 (LOC221711), mRNA
 NM_194300 Homo sapiens hypothetical protein LOC223075 (LOC223075), mRNA
 NM_194302 Homo sapiens hypothetical protein DKFZp434O0527 (DKFZp434O0527), m
 NM_194303 Homo sapiens chromosome 10 open reading frame 39 (C10orf39), mRNA
 NM_194309 Homo sapiens chromosome 21 open reading frame 125 (C21orf125), mRNA
 NM_194310 Homo sapiens hypothetical protein LOC284837 (LOC284837), mRNA
 NM_194312 Homo sapiens hypothetical protein LOC339768 (LOC339768), mRNA
 NM_194313 Homo sapiens chromosome 9 open reading frame 48 (C9orf48), mRNA
 NM_194314 Homo sapiens FRBZ1 protein (FRBZ1), mRNA
 NM_194315 Homo sapiens ubiquitin-conjugating enzyme E2, J2 (UBC8 homolog, yeast) f
 NM_194316 Homo sapiens ubiquitin-conjugating enzyme E2, J2 (UBC6 homolog, yeast) f
 NM_194317 Homo sapiens hypothetical protein MGC52057 (MGC52057), mRNA
 NM_194318 Homo sapiens beta 3-glycosyltransferase-like (B3GTL), mRNA
 NM_194319 Homo sapiens zinc finger protein 542 (ZNF542), mRNA
 NM_194320 Homo sapiens zinc finger protein 169 (ZNF169), mRNA
 NM_194322 Homo sapiens otoferlin (OTOF), transcript variant 3, mRNA
 NM_194323 Homo sapiens otoferlin (OTOF), transcript variant 4, mRNA
 NM_194324 Homo sapiens hypothetical protein MGC39900 (MGC39900), mRNA
 NM_194325 Homo sapiens zinc finger protein 30 (KOX 28) (ZNF30), mRNA
 NM_194326 Homo sapiens hypothetical protein MGC52010 (MGC52010), mRNA
 NM_194327 Homo sapiens galectin-3 internal gene (GALIG), mRNA
 NM_194328 Homo sapiens ring finger protein 38 (RNF38), transcript variant 2, mRNA
 NM_194329 Homo sapiens ring finger protein 38 (RNF38), transcript variant 3, mRNA
 NM_194330 Homo sapiens ring finger protein 38 (RNF38), transcript variant 5, mRNA
 NM_194331 Homo sapiens ring finger protein 38 (RNF38), transcript variant 4, mRNA
 NM_194332 Homo sapiens ring finger protein 38 (RNF38), transcript variant 6, mRNA
 NM_194352 Homo sapiens ring finger protein 40 (RNF40), transcript variant 2, mRNA
 NM_194356 Homo sapiens epimorphin (EPIM), transcript variant 2, mRNA
 NM_194358 Homo sapiens ring finger protein 41 (RNF41), transcript variant 2, mRNA
 NM_194359 Homo sapiens ring finger protein 41 (RNF41), transcript variant 3, mRNA
 NM_194428 Homo sapiens DEAH (Asp-Glu-Ala-His) box polypeptide 34 (DHX34), transcr
 NM_194429 Homo sapiens FGFR1 oncogene partner (FGFR1OP), transcript variant 2, m
 NM_194430 Homo sapiens ribonuclease, RNase A family, 4 (RNASE4), transcript variant
 NM_194431 Homo sapiens ribonuclease, RNase A family, 4 (RNASE4), transcript variant
 NM_194434 Homo sapiens VAMP (vesicle-associated membrane protein)-associated pro
 NM_194435 Homo sapiens vasoactive intestinal peptide (VIP), transcript variant 2, mRNA
 NM_194436 Homo sapiens lactate dehydrogenase D (LDHD), nuclear gene encoding mit
 NM_194439 Homo sapiens hypothetical protein LOC285498 (LOC285498), mRNA
 NM_194441 Homo sapiens butyrophilin, subfamily 3, member A1 (BTN3A1), transcript va

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NM_194442 Homo sapiens lamin B receptor (LBR), transcript variant 2, mRNA
 NM_194447 Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain
 NM_194448 Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain
 NM_194449 Homo sapiens pleckstrin homology domain containing, family E (with leucine
 NM_194450 Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain
 NM_194451 Homo sapiens lipoic acid synthetase (LIAS), nuclear gene encoding mitocho
 NM_194452 Homo sapiens ring finger protein 121 (RNF121), transcript variant 2, mRNA
 NM_194453 Homo sapiens ring finger protein 121 (RNF121), transcript variant 3, mRNA
 NM_194454 Homo sapiens cerebral cavernous malformations 1 (CCM1), transcript varian
 NM_194455 Homo sapiens cerebral cavernous malformations 1 (CCM1), transcript varian
 NM_194456 Homo sapiens cerebral cavernous malformations 1 (CCM1), transcript varian
 NM_194457 Homo sapiens ubiquitin-conjugating enzyme E2, J2 (UBC6 homolog, yeast) (I
 NM_194458 Homo sapiens ubiquitin-conjugating enzyme E2, J2 (UBC6 homolog, yeast) (I
 NM_194460 Homo sapiens ring finger protein 126 (RNF126), transcript variant 2, mRNA
 NM_194463 Homo sapiens ring finger protein 128 (RNF128), transcript variant 1, mRNA
 NM_197939 Homo sapiens ring finger protein 135 (RNF135), transcript variant 2, mRNA
 NM_197941 Homo sapiens similar to ADAMTS-10 precursor (A disintegrin and metallopro
 NM_197947 Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain
 NM_197948 Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain
 NM_197949 Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain
 NM_197950 Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain
 NM_197951 Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain
 NM_197952 Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain
 NM_197953 Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain
 NM_197954 Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain
 NM_197955 Homo sapiens normal mucosa of esophagus specific 1 (NMES1), transcript v
 NM_197956 Homo sapiens chromosome 9 open reading frame 90 (C9orf90), mRNA
 NM_197957 Homo sapiens MAX protein (MAX), transcript variant 6, mRNA
 NM_197958 Homo sapiens acheron (FLJ11196), transcript variant 2, mRNA
 NM_197980 Homo sapiens dipeptidylpeptidase 8 (DPP8), transcript variant 3, mRNA
 NM_197961 Homo sapiens dipeptidylpeptidase 8 (DPP8), transcript variant 4, mRNA
 NM_197962 Homo sapiens glutaredoxin 2 (GLRX2), transcript variant 2, mRNA
 NM_197964 Homo sapiens hypothetical protein HSPC268 (HSPC268), mRNA
 NM_197985 Homo sapiens sodium-dependent organic anion transporter (SOAT), mRNA
 NM_197966 Homo sapiens BH3 interacting domain death agonist (BID), transcript variant
 NM_197967 Homo sapiens BH3 interacting domain death agonist (BID), transcript variant
 NM_197988 Homo sapiens zinc finger protein 198 (ZNF198), mRNA
 NM_197970 Homo sapiens bol, boule-like (Drosophila) (BOLL), transcript variant 1, mRNA
 NM_197972 Homo sapiens non-metastatic cells 7, protein expressed in (nucleoside-diph
 NM_197973 Homo sapiens asparagine-linked glycosylation 2 homolog (yeast, alpha-1,3-r
 NM_197974 Homo sapiens butyrophilin, subfamily 3, member A3 (BTN3A3), transcript va
 NM_197975 Homo sapiens butyrophilin-like 3 (BTN3L3), transcript variant 1, mRNA
 NM_197976 Homo sapiens PBX/knotted 1 homeobox 1 (PKNOX1), transcript variant 2, m
 NM_197977 Homo sapiens zinc finger protein 189 (ZNF189), mRNA
 NM_197978 Homo sapiens hemogen (HEMGN), transcript variant 2, mRNA
 NM_198038 Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 9 (I
 NM_198039 Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 9 (I
 NM_198040 Homo sapiens polyhomeotic-like 2 (Drosophila) (PHC2), transcript variant 1,
 NM_198041 Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 6 (I
 NM_198042 Homo sapiens PDZ and LIM domain 2 (mystique) (PDLIM2), transcript varian
 NM_198043 Homo sapiens zinc finger, DHHC domain containing 16 (ZDHHC16), transcri
 NM_198044 Homo sapiens zinc finger, DHHC domain containing 16 (ZDHHC16), transcri
 NM_198045 Homo sapiens zinc finger, DHHC domain containing 16 (ZDHHC16), transcri
 NM_198046 Homo sapiens zinc finger, DHHC domain containing 16 (ZDHHC16), transcri
 NM_198047 Homo sapiens 3-hydroxyisobutyryl-Coenzyme A hydrolase (HIBCH), transcri
 NM_198053 Homo sapiens CD3Z antigen, zeta polypeptide (TIT3 complex) (CD3Z), trans
 NM_198055 Homo sapiens zinc finger protein 42 (myeloid-specific retinoic acid-responsiv

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NM_198056 Homo sapiens sodium channel, voltage-gated, type V, alpha (long QT syndr
 NM_198057 Homo sapiens delta sleep inducing peptide, immunoreactor (DSIP), transcri
 NM_198058 Homo sapiens zinc finger protein 266 (ZNF266), mRNA
 NM_198060 Homo sapiens nebulin-related anchoring protein (NRAP), transcript variant 2
 NM_198061 Homo sapiens carboxylesterase 2 (intestine, liver) (CES2), transcript variant
 NM_198066 Homo sapiens glucosamine-phosphate N-acetyltransferase 1 (GNPNAT1), tr
 NM_198074 Homo sapiens olfactory receptor, family 2, subfamily C, member 3 (OR2C3),
 NM_198075 Homo sapiens hypothetical protein DKFZp761L1518 (DKFZp761L1518), mR
 NM_198076 Homo sapiens family with sequence similarity 36, member A (FAM36A), mR
 NM_198077 Homo sapiens gmi117 (gmi117), mRNA
 NM_198078 Homo sapiens chromosome 21 open reading frame 121 (C21orf121), mRNA
 NM_198079 Homo sapiens similar to golgi autoantigen, golgin subfamily a (FLJ40113), m
 NM_198080 Homo sapiens hypothetical protein LOC253827 (LOC253827), mRNA
 NM_198081 Homo sapiens sex comb on midleg-like 4 (Drosophila) (SCML4), mRNA
 NM_198082 Homo sapiens hypothetical protein LOC284001 (LOC284001), mRNA
 NM_198083 Homo sapiens dehydrogenase/reductase (SDR family) member 4 like 2 (DHF
 NM_198085 Homo sapiens ring finger protein 148 (RNF148), mRNA
 NM_198086 Homo sapiens jub, ajuba homolog (Xenopus laevis) (JUB), transcript variant
 NM_198087 Homo sapiens zinc finger protein 200 (ZNF200), mRNA
 NM_198088 Homo sapiens zinc finger protein 200 (ZNF200), mRNA
 NM_198089 Homo sapiens zinc finger protein 155 (pH2-96) (ZNF155), transcript variant 2
 NM_198097 Homo sapiens chromosome 7 open reading frame 28B (C7orf28B), mRNA
 NM_198098 Homo sapiens aquaporin 1 (channel-forming integral protein, 28kDa) (AQP1),
 NM_198120 Homo sapiens estrogen receptor binding site associated, antigen, 9 (EBAG9
 NM_198123 Homo sapiens CUB and Sushi multiple domains 3 (CSMD3), transcript varia
 NM_198124 Homo sapiens CUB and Sushi multiple domains 3 (CSMD3), transcript varia
 NM_198125 Homo sapiens TYRO protein tyrosine kinase binding protein (TYROBP), tran
 NM_198128 Homo sapiens ring finger protein 138 (RNF138), transcript variant 2, mRNA
 NM_198129 Homo sapiens laminin, alpha 3 (LAMA3), transcript variant 1, mRNA
 NM_198138 Homo sapiens SEC31-like 2 (S. cerevisiae) (SEC31L2), transcript variant 2, i
 NM_198139 Homo sapiens semenogelin I (SEMG1), transcript variant 2, mRNA
 NM_198141 Homo sapiens glucosidase, alpha; neutral C (GANC), mRNA
 NM_198147 Homo sapiens hypothetical protein LOC116236 (LOC116236), mRNA
 NM_198148 Homo sapiens carboxypeptidase X (M14 family), member 2 (CPXM2), mRNA
 NM_198149 Homo sapiens chromosome 1 open reading frame 40 (C1orf40), mRNA
 NM_198150 Homo sapiens hypothetical protein DKFZp313G1735 (DKFZp313G1735), m
 NM_198151 Homo sapiens hypothetical protein LOC253012 (LOC253012), mRNA
 NM_198152 Homo sapiens urotensin II-related peptide (URP), mRNA
 NM_198153 Homo sapiens triggering receptor expressed on myeloid cells-like 4 (TREML
 NM_198154 Homo sapiens hypothetical protein LOC339168 (LOC339168), mRNA
 NM_198155 Homo sapiens chromosome 21 open reading frame 33 (C21orf33), nuclear g
 NM_198156 Homo sapiens von Hippel-Lindau syndrome (VHL), transcript variant 2, mRN
 NM_198157 Homo sapiens ubiquitin-conjugating enzyme E2L 3 (UBE2L3), transcript vari
 NM_198158 Homo sapiens microphthalmia-associated transcription factor (MITF), transcr
 NM_198159 Homo sapiens microphthalmia-associated transcription factor (MITF), transcr
 NM_198173 Homo sapiens transcription factor CP2-like 4 (TFCP2L4), transcript variant 2
 NM_198174 Homo sapiens transcription factor CP2-like 4 (TFCP2L4), transcript variant 3
 NM_198175 Homo sapiens non-metastatic cells 1, protein (NM23A) expressed in (NME1)
 NM_198177 Homo sapiens microphthalmia-associated transcription factor (MITF), transcr
 NM_198178 Homo sapiens microphthalmia-associated transcription factor (MITF), transcr
 NM_198179 Homo sapiens G protein-coupled receptor 103 (GPR103), mRNA
 NM_198180 Homo sapiens P518 precursor protein (P518), mRNA
 NM_198181 Homo sapiens similar to golgi autoantigen, golgin subfamily a, 2; SY11 prote
 NM_198182 Homo sapiens transcription factor CP2-like 2 (TFCP2L2), transcript variant 2
 NM_198183 Homo sapiens ubiquitin-conjugating enzyme E2L 6 (UBE2L6), transcript vari
 NM_198184 Homo sapiens osteonin (OSTN), mRNA
 NM_198185 Homo sapiens oviductin protease (OVTN), mRNA

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NM_198186 Homo sapiens astrotactin 2 (ASTN2), transcript variant 2, mRNA
 NM_198187 Homo sapiens astrotactin 2 (ASTN2), transcript variant 3, mRNA
 NM_198188 Homo sapiens astrotactin 2 (ASTN2), transcript variant 4, mRNA
 NM_198189 Homo sapiens COP9 constitutive photomorphogenic homolog subunit 8 (P
 NM_198194 Homo sapiens stomatin (STOM), transcript variant 2, mRNA
 NM_198195 Homo sapiens ubiquitin-activating enzyme E1C (UBA3 homolog, yeast) (UB
 NM_198196 Homo sapiens CD96 antigen (CD96), transcript variant 1, mRNA
 NM_198197 Homo sapiens ubiquitin-activating enzyme E1C (UBA3 homolog, yeast) (UB
 NM_198201 Homo sapiens processing of precursor 5, ribonuclease P/MRP subunit (S. ce
 NM_198202 Homo sapiens processing of precursor 5, ribonuclease P/MRP subunit (S. ce
 NM_198204 Homo sapiens transcription factor-like 4 (TCFL4), transcript variant 2, mRNA
 NM_198205 Homo sapiens transcription factor-like 4 (TCFL4), transcript variant 1, mRNA
 NM_198207 Homo sapiens LAG1 longevity assurance homolog 1 (S. cerevisiae) (LASS1)
 NM_198212 Homo sapiens caveolin 2 (CAV2), transcript variant 2, mRNA
 NM_198213 Homo sapiens 2'-5'-oligoadenylate synthetase-like (OASL), transcript variant
 NM_198215 Homo sapiens family with sequence similarity 13, member C1 (FAM13C1), tr
 NM_198216 Homo sapiens small nuclear ribonucleoprotein polypeptides B and B1 (SNRP
 NM_198217 Homo sapiens inhibitor of growth family, member 1 (ING1), transcript variant
 NM_198218 Homo sapiens inhibitor of growth family, member 1 (ING1), transcript variant
 NM_198219 Homo sapiens inhibitor of growth family, member 1 (ING1), transcript variant
 NM_198220 Homo sapiens small nuclear ribonucleoprotein polypeptide B' (SNRPB2), tra
 NM_198225 Homo sapiens Rho-related BTB domain containing 1 (RHOBTB1), transcript
 NM_198227 Homo sapiens regulator of G-protein signalling 12 (RGS12), transcript varian
 NM_198229 Homo sapiens regulator of G-protein signalling 12 (RGS12), transcript varian
 NM_198230 Homo sapiens regulator of G-protein signalling 12 (RGS12), transcript varian
 NM_198232 Homo sapiens ribonuclease, RNase A family, 1 (pancreatic) (RNASE1), tran
 NM_198234 Homo sapiens ribonuclease, RNase A family, 1 (pancreatic) (RNASE1), tran
 NM_198235 Homo sapiens ribonuclease, RNase A family, 1 (pancreatic) (RNASE1), tran
 NM_198236 Homo sapiens Rho guanine nucleotide exchange factor (GEF) 11 (ARHGEF
 NM_198239 Homo sapiens WNT1 inducible signaling pathway protein 3 (WISP3), transcr
 NM_198240 Homo sapiens retin (Reed-Steinberg cell-expressed intermediate filament-a
 NM_198241 Homo sapiens eukaryotic translation initiation factor 4 gamma, 1 (EIF4G1), t
 NM_198242 Homo sapiens eukaryotic translation initiation factor 4 gamma, 1 (EIF4G1), t
 NM_198243 Homo sapiens ankryrin repeat and SOCS box-containing 7 (ASB7), transcript
 NM_198244 Homo sapiens eukaryotic translation initiation factor 4 gamma, 1 (EIF4G1), t
 NM_198252 Homo sapiens gelsolin (amyloidosis, Finnish type) (GSN), transcript variant 2
 NM_198253 Homo sapiens telomerase reverse transcriptase (TERT), transcript variant 3,
 NM_198254 Homo sapiens telomerase reverse transcriptase (TERT), transcript variant 4,
 NM_198255 Homo sapiens telomerase reverse transcriptase (TERT), transcript variant 2,
 NM_198256 Homo sapiens E2F transcription factor 6 (E2F6), transcript variant b, mRNA
 NM_198257 Homo sapiens E2F transcription factor 6 (E2F6), transcript variant e, mRNA
 NM_198258 Homo sapiens E2F transcription factor 6 (E2F6), transcript variant d, mRNA
 NM_198261 Homo sapiens similar to splicing factor, arginine/serine-rich 4 (FLJ11021), tr
 NM_198262 Homo sapiens similar to splicing factor, arginine/serine-rich 4 (FLJ11021), tr
 NM_198263 Homo sapiens similar to splicing factor, arginine/serine-rich 4 (FLJ11021), tr
 NM_198264 Homo sapiens chromosome 1 open reading frame 2 (C1orf2), transcript varie
 NM_198265 Homo sapiens SPO11 meiotic protein covalently bound to DSB-like (S. cerev
 NM_198266 Homo sapiens inhibitor of growth family, member 3 (ING3), transcript variant
 NM_198267 Homo sapiens inhibitor of growth family, member 3 (ING3), transcript variant
 NM_198268 Homo sapiens homeodomain interacting protein kinase 1 (HIPK1), transcript
 NM_198269 Homo sapiens homeodomain interacting protein kinase 1 (HIPK1), transcript
 NM_198270 Homo sapiens Nance-Horan syndrome (congenital cataracts and dental anoi
 NM_198271 Homo sapiens leiomodlin 3 (fetal) (LMOD3), mRNA
 NM_198274 Homo sapiens SET and MYND domain containing 1 (SMYD1), mRNA
 NM_198275 Homo sapiens hypothetical protein LOC196264 (LOC196264), mRNA
 NM_198276 Homo sapiens transmembrane protein 17 (TMEM17), mRNA
 NM_198277 Homo sapiens solute carrier family 37 (glycerol-3-phosphate transporter), me

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NM_198278 Homo sapiens hypothetical protein LOC255743 (LOC255743), mRNA
 NM_198279 Homo sapiens chromosome X open reading frame 23 (CXorf23), mRNA
 NM_198281 Homo sapiens hypothetical protein LOC285513 (LOC285513), mRNA
 NM_198282 Homo sapiens hypothetical protein LOC340061 (LOC340061), mRNA
 NM_198283 Homo sapiens EGF-like-domain, multiple 11 (EGFL11), mRNA
 NM_198284 Homo sapiens hypothetical protein LOC349114 (LOC349114), mRNA
 NM_198285 Homo sapiens hypothetical protein LOC349136 (LOC349136), mRNA
 NM_198287 Homo sapiens inhibitor of growth family, member 4 (ING4), transcript variant
 NM_198289 Homo sapiens cell death-inducing DFFA-like effector a (CIDEA), transcript v
 NM_198291 Homo sapiens v-src sarcoma (Schmidt-Ruppin A-2) viral oncogene homolog
 NM_198309 Homo sapiens tetratricopeptide repeat domain 8 (TTC8), transcript variant 1,
 NM_198310 Homo sapiens tetratricopeptide repeat domain 8 (TTC8), transcript variant 2,
 NM_198312 Homo sapiens TAK1-binding protein 3 (TAB3), transcript variant 2, mRNA
 NM_198315 Homo sapiens loss of heterozygosity, 11, chromosomal region 2, gene A (LC
 NM_198316 Homo sapiens tensin like C1 domain containing phosphatase (TENC1), tran
 NM_198317 Homo sapiens kelch-like 17 (Drosophila) (KLHL17), mRNA
 NM_198318 Homo sapiens HMT1 hnRNP methyltransferase-like 2 (S. cerevisiae) (HRMT
 NM_198319 Homo sapiens HMT1 hnRNP methyltransferase-like 2 (S. cerevisiae) (HRMT
 NM_198320 Homo sapiens carboxypeptidase M (CPM), transcript variant 2, mRNA
 NM_198321 Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylga
 NM_198324 Homo sapiens citrate synthase (CS), nuclear gene encoding mitochondrial p
 NM_198325 Homo sapiens E2F transcription factor 6 (E2F6), transcript variant c, mRNA
 NM_198327 Homo sapiens suppression of tumorigenicity 7 like (ST7L), transcript variant
 NM_198328 Homo sapiens suppression of tumorigenicity 7 like (ST7L), transcript variant
 NM_198329 Homo sapiens ubiquitin-activating enzyme E1-domain containing 1 (UBE1D
 NM_198330 Homo sapiens inositol polyphosphate-5-phosphatase F (INPP5F), transcript
 NM_198331 Homo sapiens inositol polyphosphate-5-phosphatase F (INPP5F), transcript
 NM_198333 Homo sapiens purinergic receptor P2Y, G-protein coupled, 10 (P2RY10), tra
 NM_198334 Homo sapiens glucosidase, alpha; neutral AB (GANAB), mRNA
 NM_198335 Homo sapiens glucosidase, alpha; neutral AB (GANAB), mRNA
 NM_198336 Homo sapiens insulin induced gene 1 (INSIG1), transcript variant 2, mRNA
 NM_198337 Homo sapiens insulin induced gene 1 (INSIG1), transcript variant 3, mRNA
 NM_198353 Homo sapiens potassium channel tetramerisation domain containing 8 (KCT
 NM_198376 Homo sapiens calcium channel, voltage-dependent, alpha 1G subunit (CAC
 NM_198377 Homo sapiens calcium channel, voltage-dependent, alpha 1G subunit (CAC
 NM_198378 Homo sapiens calcium channel, voltage-dependent, alpha 1G subunit (CAC
 NM_198379 Homo sapiens calcium channel, voltage-dependent, alpha 1G subunit (CAC
 NM_198380 Homo sapiens calcium channel, voltage-dependent, alpha 1G subunit (CAC
 NM_198381 Homo sapiens E74-like factor 5 (ets domain transcription factor) (ELF5), tran
 NM_198382 Homo sapiens calcium channel, voltage-dependent, alpha 1G subunit (CAC
 NM_198383 Homo sapiens calcium channel, voltage-dependent, alpha 1G subunit (CAC
 NM_198384 Homo sapiens calcium channel, voltage-dependent, alpha 1G subunit (CAC
 NM_198385 Homo sapiens calcium channel, voltage-dependent, alpha 1G subunit (CAC
 NM_198386 Homo sapiens calcium channel, voltage-dependent, alpha 1G subunit (CAC
 NM_198387 Homo sapiens calcium channel, voltage-dependent, alpha 1G subunit (CAC
 NM_198388 Homo sapiens calcium channel, voltage-dependent, alpha 1G subunit (CAC
 NM_198389 Homo sapiens lung type-I cell membrane-associated glycoprotein (T1A-2), tr
 NM_198390 Homo sapiens C-Maf-inducing protein (CMIP), transcript variant C-mip, mRNA
 NM_198391 Homo sapiens fibronectin leucine rich transmembrane protein 3 (FLRT3), tra
 NM_198392 Homo sapiens transcription factor 21 (TCF21), transcript variant 1, mRNA
 NM_198393 Homo sapiens testis expressed sequence 14 (TEX14), transcript variant 1, r
 NM_198394 Homo sapiens chromosome 9 open reading frame 85 (C9orf85), transcript v
 NM_198395 Homo sapiens Ras-GTPase-activating protein SH3-domain-binding protein (t
 NM_198396 Homo sapiens calcium channel, voltage-dependent, alpha 1G subunit (CAC
 NM_198397 Homo sapiens calcium channel, voltage-dependent, alpha 1G subunit (CAC
 NM_198398 Homo sapiens serologically defined breast cancer antigen 84 (SDBCAG84),
 NM_198399 Homo sapiens cyclic AMP-regulated phosphoprotein, 21 kD (ARPP-21), tran

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NM_198400 Homo sapiens neural precursor cell expressed, developmentally down-regulated
 NM_198401 Homo sapiens hypothetical protein LOC157567 (LOC157567), mRNA
 NM_198402 Homo sapiens protein tyrosine phosphatase-like (proline instead of catalytic)
 NM_198403 Homo sapiens monocyte to macrophage differentiation-associated 2 (MMD2)
 NM_198404 Homo sapiens potassium channel tetramerisation domain containing 4 (KCTD)
 NM_198406 Homo sapiens progesterone and adiponectin receptor family member VI (PAQR6), tr
 NM_198407 Homo sapiens growth hormone secretagogue receptor (GHSR), transcript va
 NM_198426 Homo sapiens putative breast adenocarcinoma marker (32kD) (BC-2), transc
 NM_198427 Homo sapiens brevican (BCAN), transcript variant 2, mRNA
 NM_198428 Homo sapiens parathyroid hormone-responsive B1 gene (B1), transcript vari
 NM_198430 Homo sapiens regulator of G-protein signalling 12 (RGS12), transcript varian
 NM_198431 Homo sapiens heat shock 70kDa protein 4 (HSPA4), transcript variant 2, mR
 NM_198432 Homo sapiens regulator of G-protein signalling 12 (RGS12), transcript varian
 NM_198433 Homo sapiens serine/threonine kinase 6 (STK6), transcript variant 1, mRNA
 NM_198434 Homo sapiens serine/threonine kinase 6 (STK6), transcript variant 3, mRNA
 NM_198435 Homo sapiens serine/threonine kinase 6 (STK6), transcript variant 4, mRNA
 NM_198436 Homo sapiens serine/threonine kinase 6 (STK6), transcript variant 5, mRNA
 NM_198437 Homo sapiens serine/threonine kinase 6 (STK6), transcript variant 6, mRNA
 NM_198439 Homo sapiens kelch repeat and BTB (POZ) domain containing 3 (KBTBD3),
 NM_198440 Homo sapiens chromosome 22 open reading frame 14 (C22orf14), transcript
 NM_198441 Homo sapiens FLJ40296 protein (FLJ40296), mRNA
 NM_198442 Homo sapiens FLJ45651 protein (FLJ45651), mRNA
 NM_198443 Homo sapiens MRC2446 (UNQ2446), mRNA
 NM_198444 Homo sapiens GPAD9366 (UNQ9366), mRNA
 NM_198445 Homo sapiens FLJ45909 protein (FLJ45909), mRNA
 NM_198446 Homo sapiens FLJ45459 protein (FLJ45459), mRNA
 NM_198447 Homo sapiens FLJ42654 protein (FLJ42654), mRNA
 NM_198448 Homo sapiens LPPM429 (UNQ429), mRNA
 NM_198449 Homo sapiens similar to emblin (MGC71745), mRNA
 NM_198450 Homo sapiens AAIR8193 (UNQ8193), mRNA
 NM_198451 Homo sapiens forkhead box R2 (FOXR2), mRNA
 NM_198452 Homo sapiens pregnancy upregulated non-ubiquitously expressed CaM kina
 NM_198457 Homo sapiens zinc finger protein 600 (ZNF600), mRNA
 NM_198458 Homo sapiens zinc finger protein 497 (ZNF497), mRNA
 NM_198459 Homo sapiens FLJ37099 protein (FLJ37099), mRNA
 NM_198460 Homo sapiens hypothetical protein DKFZp686G0786 (DKFZp686G0786), m
 NM_198461 Homo sapiens FLJ45273 protein (FLJ45273), mRNA
 NM_198462 Homo sapiens FLJ46154 protein (FLJ46154), mRNA
 NM_198463 Homo sapiens FLJ42117 protein (FLJ42117), mRNA
 NM_198464 Homo sapiens tryptophan/serine protease (UNQ9391), mRNA
 NM_198465 Homo sapiens NIK related kinase (NRK), mRNA
 NM_198466 Homo sapiens FLJ37183 protein (FLJ37183), mRNA
 NM_198467 Homo sapiens FLJ42526 protein (FLJ42526), mRNA
 NM_198468 Homo sapiens chromosome 6 open reading frame 167 (C6orf167), mRNA
 NM_198469 Homo sapiens chromosome 9 open reading frame 18 (C9orf18), mRNA
 NM_198471 Homo sapiens FLJ46061 protein (FLJ46061), mRNA
 NM_198472 Homo sapiens chromosome 10 open reading frame 125 (C10orf125), mRNA
 NM_198473 Homo sapiens FLJ46111 protein (FLJ46111), mRNA
 NM_198474 Homo sapiens olfactomedin-like 1 (OLFM1), mRNA
 NM_198476 Homo sapiens FLJ41131 protein (FLJ41131), mRNA
 NM_198477 Homo sapiens DMC (UNQ473), mRNA
 NM_198478 Homo sapiens NTPase, KAP family P-loop domain containing 1 (NKPD1), m
 NM_198479 Homo sapiens FLJ40321 protein (FLJ40321), mRNA
 NM_198480 Homo sapiens zinc finger protein 615 (ZNF615), mRNA
 NM_198481 Homo sapiens LAIR hlog (UNQ3033), mRNA
 NM_198482 Homo sapiens similar to B-cell linker; B cell linker protein (LOC284948), tran
 NM_198483 Homo sapiens FLJ46536 protein (FLJ46536), mRNA

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NM_198484 Homo sapiens zinc finger protein 621 (ZNF621), mRNA
 NM_198485 Homo sapiens FLJ41238 protein (FLJ41238), mRNA
 NM_198486 Homo sapiens ribosomal protein L7-like 1 (RPL7L1), mRNA
 NM_198488 Homo sapiens FLJ46072 protein (FLJ46072), mRNA
 NM_198489 Homo sapiens similar to DLNB14 (DLNB14), mRNA
 NM_198490 Homo sapiens RAB43, member RAS oncogene family (RAB43), mRNA
 NM_198491 Homo sapiens FLJ44299 protein (FLJ44299), mRNA
 NM_198492 Homo sapiens liver and lymph node sinusoidal endothelial cell C-type lectin (I
 NM_198493 Homo sapiens FLJ45235 protein (FLJ45235), mRNA
 NM_198494 Homo sapiens FLJ16030 protein (FLJ16030), mRNA
 NM_198495 Homo sapiens CTAGE family, member 4 (CTAGE4), mRNA
 NM_198496 Homo sapiens A-domain containing protein similar to matrilin and collagen (I
 NM_198498 Homo sapiens Similar to RIKEN cDNA 1810046K07 gene (MGC50104), mRf
 NM_198499 Homo sapiens FLJ46156 protein (FLJ46156), mRNA
 NM_198501 Homo sapiens FLJ42461 protein (FLJ42461), mRNA
 NM_198502 Homo sapiens FLJ43826 protein (FLJ43826), mRNA
 NM_198503 Homo sapiens sodium- and chloride-activated ATP-sensitive potassium char
 NM_198504 Homo sapiens progesterin and adipoQ receptor family member IX (PAQR9), m
 NM_198506 Homo sapiens FLJ44691 protein (FLJ44691), mRNA
 NM_198507 Homo sapiens HGS_RE408 (UNQ1912), mRNA
 NM_198508 Homo sapiens FLJ44186 protein (FLJ44186), mRNA
 NM_198510 Homo sapiens ITI-like protein (UNQ6369), mRNA
 NM_198511 Homo sapiens FLJ42925 protein (FLJ42925), mRNA
 NM_198512 Homo sapiens FLJ25989 protein (FLJ25989), mRNA
 NM_198513 Homo sapiens CGI-72 protein (CGI-72), transcript variant 3, mRNA
 NM_198514 Homo sapiens NHL repeat containing 2 (NHLRC2), mRNA
 NM_198516 Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylga
 NM_198517 Homo sapiens FLJ00332 protein (FLJ00332), mRNA
 NM_198519 Homo sapiens similar to Cytochrome c, somatic (MGC12965), mRNA
 NM_198520 Homo sapiens FLJ44112 protein (FLJ44112), mRNA
 NM_198521 Homo sapiens FLJ25323 protein (FLJ25323), mRNA
 NM_198524 Homo sapiens LOC161577 protein (LOC161577), mRNA
 NM_198525 Homo sapiens similar to kinesin family member 21A; N-5 kinesin (LOC37465
 NM_198526 Homo sapiens hypothetical protein DKFZp547K1113 (DKFZp547K1113), mF
 NM_198527 Homo sapiens Similar to RIKEN cDNA 1110033O09 gene (MGC45386), mRf
 NM_198529 Homo sapiens FLJ46247 protein (FLJ46247), mRNA
 NM_198531 Homo sapiens ATPase, Class II, type 9B (ATP9B), mRNA
 NM_198532 Homo sapiens chromosome 19 open reading frame 35 (C19orf35), mRNA
 NM_198533 Homo sapiens short-chain dehydrogenase/reductase 10 (SCDR10), transcript
 NM_198534 Homo sapiens FLJ35784 protein (FLJ35784), mRNA
 NM_198537 Homo sapiens FLJ44968 protein (FLJ44968), mRNA
 NM_198538 Homo sapiens HJAR698 (UNQ698), mRNA
 NM_198539 Homo sapiens zinc finger protein 568 (ZNF568), mRNA
 NM_198540 Homo sapiens UDP-Gal:betaGal beta 1,3-galactosyltransferase polypeptide
 NM_198541 Homo sapiens insulin growth factor-like family member 1 (IGFL1), mRNA
 NM_198542 Homo sapiens similar to hypothetical protein FLJ23233 (MGC4728), mRNA
 NM_198543 Homo sapiens LOC148872 protein (LOC148872), mRNA
 NM_198544 Homo sapiens cortistatin (CORT), transcript variant 1, mRNA
 NM_198545 Homo sapiens hypothetical gene supported by AK075558; BC021286 (LOC3
 NM_198546 Homo sapiens hypothetical protein LOC374955 (LOC374955), mRNA
 NM_198547 Homo sapiens FLJ46354 protein (FLJ46354), mRNA
 NM_198549 Homo sapiens FLJ35093 protein (FLJ35093), mRNA
 NM_198550 Homo sapiens FLJ36760 protein (FLJ36760), mRNA
 NM_198552 Homo sapiens hypothetical gene supported by BC009447 (MGC15887), mRf
 NM_198553 Homo sapiens FLJ30851 protein (FLJ30851), mRNA
 NM_198554 Homo sapiens thyroid adenoma associated (THADA), mRNA
 NM_198557 Homo sapiens FLJ45645 protein (FLJ45645), mRNA

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NM_198559 Homo sapiens hypothetical gene supported by BC052750 (MGC50811), mRNA
 NM_198560 Homo sapiens Iipoma HMGC fusion partner-like protein 4 (LOC375323), mRNA
 NM_198562 Homo sapiens FLJ43654 protein (FLJ43654), mRNA
 NM_198563 Homo sapiens Similar to RIKEN cDNA 1810038N08 gene (MGC50222), mRNA
 NM_198564 Homo sapiens FLJ44290 protein (FLJ44290), mRNA
 NM_198565 Homo sapiens ELLP3030 (UNQ3030), mRNA
 NM_198566 Homo sapiens FLJ32363 protein (FLJ32363), mRNA
 NM_198567 Homo sapiens FLJ44216 protein (FLJ44216), mRNA
 NM_198568 Homo sapiens gap junction protein, beta 7 (GJB7), mRNA
 NM_198569 Homo sapiens G protein-coupled receptor 126 (GPR126), mRNA
 NM_198570 Homo sapiens PSST739 (UNQ739), mRNA
 NM_198571 Homo sapiens FLJ39237 protein (FLJ39237), mRNA
 NM_198572 Homo sapiens similar to Putative protein C21orf56 (MGC81633), mRNA
 NM_198573 Homo sapiens GAAI470 (UNQ470), mRNA
 NM_198577 Homo sapiens FLJ46361 protein (FLJ46361), mRNA
 NM_198580 Homo sapiens solute carrier family 27 (fatty acid transporter), member 1 (SLC27A1), mRNA
 NM_198581 Homo sapiens zinc finger CCHC type domain containing 6 (ZC3HDC6), mRNA
 NM_198582 Homo sapiens FLJ43374 protein (FLJ43374), mRNA
 NM_198584 Homo sapiens carbonic anhydrase XIII (CA13), mRNA
 NM_198585 Homo sapiens GLSR2492 (UNQ2492), mRNA
 NM_198586 Homo sapiens NHL repeat containing 1 (NHLRC1), mRNA
 NM_198587 Homo sapiens regulator of G-protein signalling 12 (RGS12), transcript variant 1, mRNA
 NM_198589 Homo sapiens basigin (OK blood group) (BSG), transcript variant 2, mRNA
 NM_198590 Homo sapiens basigin (OK blood group) (BSG), transcript variant 3, mRNA
 NM_198591 Homo sapiens basigin (OK blood group) (BSG), transcript variant 4, mRNA
 NM_198593 Homo sapiens C1q and tumor necrosis factor related protein 1 (C1QTNF1), transcript variant 1, mRNA
 NM_198594 Homo sapiens C1q and tumor necrosis factor related protein 1 (C1QTNF1), transcript variant 2, mRNA
 NM_198595 Homo sapiens actin filament associated protein (AFAP), transcript variant 2, mRNA
 NM_198596 Homo sapiens sulfatase 2 (SULF2), transcript variant 2, mRNA
 NM_198597 Homo sapiens SEC24 related gene family, member C (S. cerevisiae) (SEC24), mRNA
 NM_198679 Homo sapiens Rap guanine nucleotide exchange factor (GEF) 1 (RAPGEF1), mRNA
 NM_198681 Homo sapiens putative NFkB activating protein (KIAA0720), transcript variant 1, mRNA
 NM_198682 Homo sapiens glycoporphin E (GYPE), transcript variant 2, mRNA
 NM_198686 Homo sapiens RAB15, member RAS oncogene family (RAB15), mRNA
 NM_198687 Homo sapiens keratin associated protein 10-4 (KRTAP10-4), mRNA
 NM_198688 Homo sapiens keratin associated protein 10-6 (KRTAP10-6), mRNA
 NM_198689 Homo sapiens keratin associated protein 10-7 (KRTAP10-7), mRNA
 NM_198690 Homo sapiens keratin associated protein 10-9 (KRTAP10-9), mRNA
 NM_198691 Homo sapiens keratin associated protein 10-1 (KRTAP10-1), mRNA
 NM_198692 Homo sapiens keratin associated protein 10-11 (KRTAP10-11), mRNA
 NM_198693 Homo sapiens keratin associated protein 10-2 (KRTAP10-2), mRNA
 NM_198694 Homo sapiens keratin associated protein 10-5 (KRTAP10-5), mRNA
 NM_198695 Homo sapiens keratin associated protein 10-8 (KRTAP10-8), mRNA
 NM_198696 Homo sapiens keratin associated protein 10-3 (KRTAP10-3), mRNA
 NM_198697 Homo sapiens keratin associated protein 12-3 (KRTAP12-3), mRNA
 NM_198698 Homo sapiens keratin associated protein 12-4 (KRTAP12-4), mRNA
 NM_198699 Homo sapiens keratin associated protein 10-12 (KRTAP10-12), mRNA
 NM_198700 Homo sapiens CUG triplet repeat, RNA binding protein 1 (CUGBP1), transcript variant 1, mRNA
 NM_198704 Homo sapiens short-chain dehydrogenase/reductase 10 (SCDR10), transcript variant 1, mRNA
 NM_198705 Homo sapiens short-chain dehydrogenase/reductase 10 (SCDR10), transcript variant 2, mRNA
 NM_198706 Homo sapiens short-chain dehydrogenase/reductase 10 (SCDR10), transcript variant 3, mRNA
 NM_198707 Homo sapiens short-chain dehydrogenase/reductase 10 (SCDR10), transcript variant 4, mRNA
 NM_198708 Homo sapiens short-chain dehydrogenase/reductase 10 (SCDR10), transcript variant 5, mRNA
 NM_198709 Homo sapiens arylsulfatase B (ARSB), transcript variant 2, mRNA
 NM_198712 Homo sapiens prostaglandin E receptor 3 (subtype EP3) (PTGER3), transcript variant 1, mRNA
 NM_198713 Homo sapiens prostaglandin E receptor 3 (subtype EP3) (PTGER3), transcript variant 2, mRNA
 NM_198714 Homo sapiens prostaglandin E receptor 3 (subtype EP3) (PTGER3), transcript variant 3, mRNA

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NM_198715 Homo sapiens prostaglandin E receptor 3 (subtype EP3) (PTGER3), transcrip
 NM_198716 Homo sapiens prostaglandin E receptor 3 (subtype EP3) (PTGER3), transcrip
 NM_198717 Homo sapiens prostaglandin E receptor 3 (subtype EP3) (PTGER3), transcrip
 NM_198718 Homo sapiens prostaglandin E receptor 3 (subtype EP3) (PTGER3), transcrip
 NM_198719 Homo sapiens prostaglandin E receptor 3 (subtype EP3) (PTGER3), transcrip
 NM_198720 Homo sapiens prostaglandin E receptor 3 (subtype EP3) (PTGER3), transcrip
 NM_198721 Homo sapiens collagen, type XXV, alpha 1 (COL25A1), transcript variant 1, r
 NM_198722 Homo sapiens amphoterin-induced gene and ORF 3 (AMIGO3), mRNA
 NM_198723 Homo sapiens transcription elongation factor A (SII), 2 (TCEA2), transcript va
 NM_198793 Homo sapiens CD47 antigen (RH-related antigen, integrin-associated signal
 NM_198794 Homo sapiens mitogen-activated protein kinase kinase kinase 5 (MAF
 NM_198795 Homo sapiens tudor domain containing 1 (TDRD1), mRNA
 NM_198797 Homo sapiens prostaglandin E synthase (PTGES), transcript variant 2, mRN
 NM_198798 Homo sapiens ankyrin repeat domain 5 (ANKRD5), transcript variant 2, mRN
 NM_198799 Homo sapiens breast carcinoma amplified sequence 4 (BCAS4), transcript v
 NM_198822 Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex, su
 NM_198827 Homo sapiens G protein-coupled receptor 133 (GPR133), mRNA
 NM_198828 Homo sapiens similar to microtubule associated testis specific serine/threoni
 NM_198829 Homo sapiens ras-related C3 botulinum toxin substrate 1 (rho family, small G
 NM_198830 Homo sapiens ATP citrate lyase (ACLY), transcript variant 2, mRNA
 NM_198833 Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin),
 NM_198834 Homo sapiens acetyl-Coenzyme A carboxylase alpha (ACACA), transcript va
 NM_198835 Homo sapiens acetyl-Coenzyme A carboxylase alpha (ACACA), transcript va
 NM_198836 Homo sapiens acetyl-Coenzyme A carboxylase alpha (ACACA), transcript va
 NM_198837 Homo sapiens acetyl-Coenzyme A carboxylase alpha (ACACA), transcript va
 NM_198838 Homo sapiens acetyl-Coenzyme A carboxylase alpha (ACACA), transcript va
 NM_198839 Homo sapiens acetyl-Coenzyme A carboxylase alpha (ACACA), transcript va
 NM_198841 Homo sapiens chromosome 9 open reading frame 10 opposite strand (C9orf
 NM_198843 Homo sapiens surfactant, pulmonary-associated protein B (SFTPB), transcrip
 NM_198844 Homo sapiens zona pellucida binding protein 2 (ZBP2), transcript variant 1,
 NM_198845 Homo sapiens sialic acid binding Ig-like lectin 6 (SIGLEC6), transcript varian
 NM_198846 Homo sapiens sialic acid binding Ig-like lectin 6 (SIGLEC6), transcript variarr
 NM_198847 Homo sapiens FLJ22794 protein (FLJ22794), transcript variant 2, mRNA
 NM_198849 Homo sapiens similar to seven in absentia 2 (LOC283514), mRNA
 NM_198850 Homo sapiens pleckstrin homology-like domain, family B, member 3 (PHLDE
 NM_198851 Homo sapiens hypothetical protein LOC348645 (LOC348645), mRNA
 NM_198853 Homo sapiens tripartite motif-containing 50C (TRIM50C), mRNA
 NM_198855 Homo sapiens zinc finger protein 211 (ZNF211), transcript variant 2, mRNA
 NM_198856 Homo sapiens CAP-binding protein complex interacting protein 1 (FLJ23588)
 NM_198857 Homo sapiens similar to sodium- and chloride-dependent creatine transport
 NM_198859 Homo sapiens prickle-like 2 (Drosophila) (PRICKLE2), mRNA
 NM_198867 Homo sapiens hypothetical protein MGC15677 (MGC15677), transcript varia
 NM_198868 Homo sapiens KIAA0676 protein (KIAA0676), transcript variant 1, mRNA
 NM_198880 Homo sapiens FLJ20259 protein (FLJ20259), transcript variant 2, mRNA
 NM_198881 Homo sapiens FLJ20298 protein (FLJ20298), transcript variant 2, mRNA
 NM_198883 Homo sapiens metaxin 1 (MTX1), transcript variant 2, mRNA
 NM_198887 Homo sapiens nucleoporin 43kDa (NUP43), transcript variant 1, mRNA
 NM_198889 Homo sapiens ankyrin repeat domain 17 (ANKRD17), transcript variant 2, ml
 NM_198890 Homo sapiens APG16 autophagy 16-like (S. cerevisiae) (APG16L), transcrip
 NM_198892 Homo sapiens BMP2 inducible kinase (BMP2IK), transcript variant 1, mRNA
 NM_198893 Homo sapiens zinc finger protein 160 (ZNF160), transcript variant 2, mRNA
 NM_198896 Homo sapiens RAB6A, member RAS oncogene family (RAB6A), transcript v
 NM_198897 Homo sapiens fibroblast growth factor (acidic) intracellular binding protein (F
 NM_198900 Homo sapiens fomin-like 3 (FMNL3), transcript variant 2, mRNA
 NM_198901 Homo sapiens sorcin (SRI), transcript variant 2, mRNA
 NM_198902 Homo sapiens transmembrane 4 superfamily member 8 (TM4SF8), transcript
 NM_198903 Homo sapiens gamma-aminobutyric acid (GABA) A receptor, gamma 2 (GAE

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NM_198904 Homo sapiens gamma-aminobutyric acid (GABA) A receptor, gamma 2 (GABRG2), mRNA
 NM_198907 Homo sapiens hypothetical protein LOC253982 (LOC253982), mRNA
 NM_198920 Homo sapiens chromosome 6 open reading frame 157 (C6orf157), mRNA
 NM_198923 Homo sapiens MAS-related GPR, member D (MRGPRD), mRNA
 NM_198924 Homo sapiens hypothetical protein MGC45477 (MGC45477), mRNA
 NM_198925 Homo sapiens sema domain, immunoglobulin domain (Ig), transmembrane c
 NM_198926 Homo sapiens hypothetical protein LOC55924 (LOC55924), transcript varian
 NM_198928 Homo sapiens interferon-inducible protein X (IFIX), transcript variant a2, mR
 NM_198929 Homo sapiens interferon-inducible protein X (IFIX), transcript variant b1, mR
 NM_198930 Homo sapiens interferon-inducible protein X (IFIX), transcript variant b2, mR
 NM_198935 Homo sapiens synovial sarcoma translocation gene on chromosome 18-like
 NM_198938 Homo sapiens prostaglandin E synthase 2 (PTGES2), transcript variant 2, m
 NM_198939 Homo sapiens prostaglandin E synthase 2 (PTGES2), transcript variant 3, m
 NM_198940 Homo sapiens prostaglandin E synthase 2 (PTGES2), transcript variant 4, m
 NM_198941 Homo sapiens tumor differentially expressed 1 (TDE1), transcript variant 2, n
 NM_198943 Homo sapiens CXYorf1-related protein (MGC52000), mRNA
 NM_198944 Homo sapiens olfactory receptor, family 7, subfamily C, member 1 (OR7C1),
 NM_198945 Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region,
 NM_198946 Homo sapiens lipocalin 6 (LCN6), mRNA
 NM_198947 Homo sapiens cancer-associated nucleoprotein (CANP), mRNA
 NM_198948 Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 1 (I
 NM_198949 Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 1 (I
 NM_198950 Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 1 (I
 NM_198951 Homo sapiens transglutaminase 2 (C polypeptide, protein-glutamine-gamma
 NM_198952 Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 1 (I
 NM_198953 Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 1 (I
 NM_198954 Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 1 (I
 NM_198955 Homo sapiens mannosyl (alpha-1,6)-glycoprotein beta-1,6-N-acetyl-glucosa
 NM_198956 Homo sapiens Sp8 transcription factor (SP8), transcript variant 2, mRNA
 NM_198963 Homo sapiens DEAH (Asp-Glu-Ala-Asp/His) box polypeptide 57 (DHX57), tra
 NM_198964 Homo sapiens parathyroid hormone-like hormone (PTH1L), transcript varian
 NM_198965 Homo sapiens parathyroid hormone-like hormone (PTH1L), transcript varian
 NM_198966 Homo sapiens parathyroid hormone-like hormone (PTH1L), transcript varian
 NM_198968 Homo sapiens DAZ interacting protein 1 (DZIP1), mRNA
 NM_198969 Homo sapiens amino-terminal enhancer of split (AES), transcript variant 1, r
 NM_198970 Homo sapiens amino-terminal enhancer of split (AES), transcript variant 3, r
 NM_198971 Homo sapiens MBD2 (methyl-CpG-binding protein)-interacting zinc finger pr
 NM_198973 Homo sapiens MAP kinase interacting serine/threonine kinase 1 (MKNK1), r
 NM_198974 Homo sapiens PTK9 protein tyrosine kinase 9 (PTK9), transcript variant 2, m
 NM_198976 Homo sapiens TH1-like (Drosophila) (TH1L), transcript variant 1, mRNA
 NM_198977 Homo sapiens Rho guanine nucleotide exchange factor (GEF) 1 (ARHGEF1),
 NM_198988 Homo sapiens leukocyte receptor cluster (LRC) member 9 (LENG9), mRNA
 NM_198989 Homo sapiens deleted in lymphocytic leukemia 7 (DLEU7), mRNA
 NM_198990 Homo sapiens N-acyl-phosphatidylethanolamine-hydrolyzing phospholipase
 NM_198991 Homo sapiens potassium channel tetramerisation domain containing 1 (KCT
 NM_198992 Homo sapiens synaptotagmin X (SYT10), mRNA
 NM_198993 Homo sapiens SH3 and cysteine rich domain 2 (STAC2), mRNA
 NM_198994 Homo sapiens transglutaminase 6 (TM6), mRNA
 NM_198995 Homo sapiens chromosome 18 open reading frame 34 (C18orf34), mRNA
 NM_198996 Homo sapiens membrane-associated phospholipase A1 beta (LOC375108),
 NM_198998 Homo sapiens aquaporin 12 (AQP12), mRNA
 NM_198999 Homo sapiens prestin (motor protein) (PRES), transcript variant a, mRNA
 NM_199000 Homo sapiens lipoma HMGIC fusion partner-like 3 (LHFPL3), mRNA
 NM_199001 Homo sapiens Similar to RIKEN cDNA 2310002J15 gene (MGC59937), mR
 NM_199002 Homo sapiens Rho guanine nucleotide exchange factor (GEF) 1 (ARHGEF1),
 NM_199003 Homo sapiens THAP domain containing, apoptosis associated protein 1 (TH
 NM_199004 Homo sapiens arrestin, beta 2 (ARRB2), transcript variant 2, mRNA

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NM_199005 Homo sapiens zinc finger protein 322B (ZNF322B), mRNA
 NM_199006 Homo sapiens cortistatin (CORT), transcript variant 3, mRNA
 NM_199037 Homo sapiens sodium channel, voltage-gated, type I, beta (SCN1B), transcript variant 1, mRNA
 NM_199039 Homo sapiens kelch-like 5 (Drosophila) (KLHL5), transcript variant b, mRNA
 NM_199040 Homo sapiens nudix (nu cleoside diphosphate linked moiety X)-type motif 4 (NM_199043)
 NM_199043 Homo sapiens chromosome 14 open reading frame 102 (C14orf102), transcript variant 1, mRNA
 NM_199044 Homo sapiens MGC22960 gene (MGC22960), mRNA
 NM_199046 Homo sapiens testis/prostate/placenta-expressed protein (TEPP), transcript variant 1, mRNA
 NM_199047 Homo sapiens TATA box binding protein like 2 (TBPL2), mRNA
 NM_199050 Homo sapiens chromosome 21 open reading frame 25 (C21orf25), mRNA
 NM_199051 Homo sapiens DBCCR1-like (DBCCR1L), mRNA
 NM_199052 Homo sapiens chromosome 20 open reading frame 7 (C20orf7), transcript variant 1, mRNA
 NM_199053 Homo sapiens FLJ12716 protein (FLJ12716), transcript variant 2, mRNA
 NM_199054 Homo sapiens MAP kinase-interacting serine/threonine kinase 2 (MKNK2), transcript variant 1, mRNA
 NM_199069 Homo sapiens nuclear protein E3-3 (DKFZP564J0123), transcript variant 1, mRNA
 NM_199070 Homo sapiens nuclear protein E3-3 (DKFZP564J0123), transcript variant 2, mRNA
 NM_199071 Homo sapiens chromosome 21 open reading frame 58 (C21orf58), transcript variant 1, mRNA
 NM_199072 Homo sapiens I-mfa domain-containing protein (HIC), mRNA
 NM_199073 Homo sapiens nuclear protein E3-3 (DKFZP564J0123), transcript variant 3, mRNA
 NM_199074 Homo sapiens nuclear protein E3-3 (DKFZP564J0123), transcript variant 4, mRNA
 NM_199075 Homo sapiens CBF1 interacting corepressor (CIR), transcript variant 2, mRNA
 NM_199076 Homo sapiens cyclin M2 (CNNM2), transcript variant 2, mRNA
 NM_199077 Homo sapiens cyclin M2 (CNNM2), transcript variant 3, mRNA
 NM_199078 Homo sapiens cyclin M3 (CNNM3), transcript variant 2, mRNA
 NM_199121 Homo sapiens von Willebrand factor A domain-related protein (WARP), transcript variant 1, mRNA
 NM_199122 Homo sapiens transforming growth factor beta regulator 4 (TBRG4), transcript variant 1, mRNA
 NM_199123 Homo sapiens chromosome 14 open reading frame 154 (C14orf154), transcript variant 1, mRNA
 NM_199124 Homo sapiens hypothetical protein FLJ23554 (FLJ23554), transcript variant 1, mRNA
 NM_199126 Homo sapiens zinc finger protein 585A (ZNF585A), transcript variant 2, mRNA
 NM_199127 Homo sapiens gamma-glutamyltransferase-like 4 (GGTL4), transcript variant 1, mRNA
 NM_199129 Homo sapiens ubiquitin-conjugating enzyme variant Kua (Kua), mRNA
 NM_199131 Homo sapiens ventral anterior homeobox 1 (VAX1), mRNA
 NM_199133 Homo sapiens hypothetical protein LOC134145 (LOC134145), mRNA
 NM_199135 Homo sapiens FOXD4-like 2 (FOXD4L2), mRNA
 NM_199136 Homo sapiens hypothetical protein MGC72075 (MGC72075), mRNA
 NM_199138 Homo sapiens hypothetical protein FLJ25477 (FLJ25477), transcript variant 1, mRNA
 NM_199139 Homo sapiens XIAP associated factor-1 (HSXIAPAF1), transcript variant 2, mRNA
 NM_199141 Homo sapiens coactivator-associated arginine methyltransferase 1 (CARM1), transcript variant 1, mRNA
 NM_199144 Homo sapiens ubiquitin-conjugating enzyme E2 variant 1 (UBE2V1), transcript variant 1, mRNA
 NM_199160 Homo sapiens LIM homeobox 6 (LHX6), transcript variant 2, mRNA
 NM_199161 Homo sapiens serum amyloid A1 (SAA1), transcript variant 2, mRNA
 NM_199162 Homo sapiens ADP-ribosylhydrolase like 1 (ADPRHL1), transcript variant 2, mRNA
 NM_199163 Homo sapiens CXorf1-related protein (FLJ25222), mRNA
 NM_199165 Homo sapiens adenylosuccinate synthase like 1 (ADSSL1), transcript variant 1, mRNA
 NM_199166 Homo sapiens aminolevulinic acid, delta-, synthase 1 (ALAS1), transcript variant 1, mRNA
 NM_199167 Homo sapiens clusterin-like 1 (retinal) (CLUL1), transcript variant 2, mRNA
 NM_199168 Homo sapiens chemokine (C-X-C motif) ligand 12 (stromal cell-derived factor 1) (SDF1), transcript variant 1, mRNA
 NM_199169 Homo sapiens transmembrane, prostate androgen induced RNA (TMEPA1), transcript variant 1, mRNA
 NM_199170 Homo sapiens transmembrane, prostate androgen induced RNA (TMEPA1), transcript variant 2, mRNA
 NM_199171 Homo sapiens transmembrane, prostate androgen induced RNA (TMEPA1), transcript variant 3, mRNA
 NM_199173 Homo sapiens bone gamma-carboxyglutamate (gla) protein (osteocalcin) (BGLAP), transcript variant 1, mRNA
 NM_199175 Homo sapiens chromosome 21 open reading frame 123 (C21orf123), mRNA
 NM_199176 Homo sapiens mitochondrial ribosome recycling factor (MRRF), transcript variant 1, mRNA
 NM_199177 Homo sapiens mitochondrial ribosome recycling factor (MRRF), transcript variant 2, mRNA
 NM_199179 Homo sapiens kin of IRRE like 2 (Drosophila) (KIRREL2), transcript variant 2, mRNA
 NM_199180 Homo sapiens kin of IRRE like 2 (Drosophila) (KIRREL2), transcript variant 3, mRNA
 NM_199181 Homo sapiens FLJ44670 protein (FLJ44670), mRNA

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NM_199182 Homo sapiens hLAT1-3TM (IMAA), mRNA
 NM_199183 Homo sapiens testis serine protease 5 (TESSP5), mRNA
 NM_199184 Homo sapiens chromosome 6 open reading frame 108 (C6orf108), transcript
 NM_199185 Homo sapiens nucleophosmin (nucleolar phosphoprotein B23, numatrin) (NF
 NM_199186 Homo sapiens 2,3-bisphosphoglycerate mutase (BPGM), transcript variant 2
 NM_199187 Homo sapiens keratin 18 (KRT18), transcript variant 2, mRNA
 NM_199188 Homo sapiens c-Mpl binding protein (LOC113251), transcript variant 2, mRN
 NM_199190 Homo sapiens c-Mpl binding protein (LOC113251), transcript variant 3, mRN
 NM_199191 Homo sapiens brain and reproductive organ-expressed (TNFRSF1A modulai
 NM_199192 Homo sapiens brain and reproductive organ-expressed (TNFRSF1A modulai
 NM_199193 Homo sapiens brain and reproductive organ-expressed (TNFRSF1A modulai
 NM_199194 Homo sapiens brain and reproductive organ-expressed (TNFRSF1A modulai
 NM_199202 Homo sapiens Theg homolog (mouse) (THEG), transcript variant 2, mRNA
 NM_199203 Homo sapiens ubiquitin-conjugating enzyme E2 variant 1 (Kua-UEV), transcr
 NM_199204 Homo sapiens dehydrogenase/reductase (SDR family) member 9 (DHR59),
 NM_199205 Homo sapiens p30 DBC protein (DBC-1), transcript variant 2, mRNA
 NM_199206 Homo sapiens T-cell leukemia/lymphoma 1B (TCL1B), transcript variant 2, m
 NM_199227 Homo sapiens methionine aminopeptidase 1D (MAP1D), mRNA
 NM_199228 Homo sapiens thrombopoietin (myeloproliferative leukemia virus oncogene II
 NM_199229 Homo sapiens ribulose-5-phosphate-3-epimerase (RPE), transcript variant 1.
 NM_199231 Homo sapiens glial cell derived neurotrophic factor (GDNF), transcript varian
 NM_199232 Homo sapiens allantoicase (ALLC), transcript variant 2, mRNA
 NM_199234 Homo sapiens glial cell derived neurotrophic factor (GDNF), transcript varian
 NM_199235 Homo sapiens collectin sub-family member 11 (COLEC11), transcript variant
 NM_199242 Homo sapiens unc-13 homolog D (C. elegans) (UNC13D), mRNA
 NM_199243 Homo sapiens G protein-coupled receptor 150 (GPR150), mRNA
 NM_199244 Homo sapiens forkhead box protein D4b (FOXO4b), mRNA
 NM_199245 Homo sapiens vesicle-associated membrane protein 1 (synaptobrevin 1) (VP
 NM_199246 Homo sapiens cyclin G1 (CCNG1), transcript variant 2, mRNA
 NM_199247 Homo sapiens calcium channel, voltage-dependent, beta 1 subunit (CACNB
 NM_199248 Homo sapiens calcium channel, voltage-dependent, beta 1 subunit (CACNB
 NM_199249 Homo sapiens multidrug resistance-related protein (MGC13170), mRNA
 NM_199250 Homo sapiens multidrug resistance-related protein (MGC13170), mRNA
 NM_199254 Homo sapiens transmembrane phosphoinositide 3-phosphatase and tensin I
 NM_199255 Homo sapiens transmembrane phosphoinositide 3-phosphatase and tensin I
 NM_199259 Homo sapiens transmembrane phosphatase with tensin homology (TPTE), tr
 NM_199260 Homo sapiens transmembrane phosphatase with tensin homology (TPTE), tr
 NM_199261 Homo sapiens transmembrane phosphatase with tensin homology (TPTE), tr
 NM_199262 Homo sapiens Sp6 transcription factor (SP6), mRNA
 NM_199263 Homo sapiens thrombospondin, type I, domain containing 1 (THSD1), transc
 NM_199265 Homo sapiens thrombospondin, type I, domain containing 3 (THSD3), transc
 NM_199280 Homo sapiens similar to RIKEN cDNA 4632412N22 gene (LOC165186), mR
 NM_199282 Homo sapiens likely ortholog of rat CIN5-associated multi-domain containi
 NM_199285 Homo sapiens hypothetical LOC284338 (MGC70924), mRNA
 NM_199286 Homo sapiens STELLA mRNA (LOC338759), mRNA
 NM_199290 Homo sapiens alpha-NAC protein (MGC71999), mRNA
 NM_199292 Homo sapiens tyrosine hydroxylase (TH), transcript variant 1, mRNA
 NM_199293 Homo sapiens tyrosine hydroxylase (TH), transcript variant 3, mRNA
 NM_199294 Homo sapiens cortistatin (CORT), transcript variant 4, mRNA
 NM_199295 Homo sapiens cortistatin (CORT), transcript variant 5, mRNA
 NM_199296 Homo sapiens thrombospondin, type I, domain containing 3 (THSD3), transc
 NM_199297 Homo sapiens thymocyte protein thy28 (THY28), transcript variant 2, mRNA
 NM_199298 Homo sapiens thymocyte protein thy28 (THY28), transcript variant 3, mRNA
 NM_199320 Homo sapiens PHD protein Jade-1 (JADE1), transcript variant L, mRNA
 NM_199321 Homo sapiens zona pellucida binding protein 2 (ZBP2), transcript variant 2,
 NM_199324 Homo sapiens HIV-1 induced protein HIN-1 (HSHIN1), transcript variant 1, m
 NM_199326 Homo sapiens protein phosphatase 2A 48 kDa regulatory subunit (PR48), tr

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NM_199327 Homo sapiens sprouty homolog 1, antagonist of FGF signaling (Drosophila) l
 NM_199328 Homo sapiens claudin 8 (CLDN8), mRNA
 NM_199329 Homo sapiens solute carrier family 43, member 3 (SLC43A3), mRNA
 NM_199330 Homo sapiens homer homolog 2 (Drosophila) (HOMER2), transcript variant 2
 NM_199331 Homo sapiens homer homolog 2 (Drosophila) (HOMER2), transcript variant 3
 NM_199332 Homo sapiens homer homolog 2 (Drosophila) (HOMER2), transcript variant 4
 NM_199334 Homo sapiens thyroid hormone receptor, alpha (erythroblastic leukemia viral
 NM_199335 Homo sapiens FYN binding protein (FYB-120/130) (FYB), mRNA
 NM_199336 Homo sapiens hypothetical protein DKFZp434N062 (DKFZp434N062), mRNA
 NM_199337 Homo sapiens similar to RIKEN cDNA 1810059G22 (LOC374395), mRNA
 NM_199338 Homo sapiens FLJ35171 protein (FLJ35171), mRNA
 NM_199339 Homo sapiens hypothetical protein LOC374768 (LOC374768), mRNA
 NM_199341 Homo sapiens hypothetical protein LOC374920 (LOC374920), mRNA
 NM_199342 Homo sapiens hypothetical protein LOC374968 (LOC374969), mRNA
 NM_199343 Homo sapiens FLJ90637 protein (FLJ90637), mRNA
 NM_199344 Homo sapiens hypothetical protein LOC375035 (LOC375035), mRNA
 NM_199345 Homo sapiens similar to phosphatidylinositol 4-kinase alpha (LOC375133), n
 NM_199346 Homo sapiens profilin family, member 4 (PFN4), mRNA
 NM_199348 Homo sapiens protocadherin protein CDHJ (CDHJ), transcript variant 2, mRt
 NM_199349 Homo sapiens kliein-like (LOC375616), mRNA
 NM_199350 Homo sapiens hypothetical protein LOC375759 (LOC375759), mRNA
 NM_199351 Homo sapiens chromosome 1 open reading frame 32 (C1orf32), mRNA
 NM_199352 Homo sapiens putative UST1-like organic anion transporter (LOC387601), m
 NM_199353 Homo sapiens proline-rich protein BatN1 subfamily 1 (PRB1), transcript varia
 NM_199354 Homo sapiens proline-rich protein BatN1 subfamily 1 (PRB1), transcript varia
 NM_199355 Homo sapiens a disintegrin-like and metalloprotease (repolyisin type) with th
 NM_199356 Homo sapiens thrombopoietin (myeloproliferative leukemia virus oncogene li
 NM_199357 Homo sapiens similar to human GTPase-activating protein (ARHGAP11A), n
 NM_199358 Homo sapiens forkhead box D4 like 3 (FOXO4L3), mRNA
 NM_199359 Homo sapiens tumor protein D52-like 2 (TPD52L2), transcript variant 6, mR
 NM_199360 Homo sapiens tumor protein D52-like 2 (TPD52L2), transcript variant 1, mR
 NM_199361 Homo sapiens tumor protein D52-like 2 (TPD52L2), transcript variant 2, mR
 NM_199362 Homo sapiens tumor protein D52-like 2 (TPD52L2), transcript variant 3, mR
 NM_199363 Homo sapiens tumor protein D52-like 2 (TPD52L2), transcript variant 4, mR
 NM_199367 Homo sapiens spastic paraplegia 7, paraplegin (pure and complicated autos
 NM_199368 Homo sapiens transient receptor potential cation channel, subfamily C, mem
 NM_199413 Homo sapiens protein regulator of cytokinesis 1 (PRC1), transcript variant 2,
 NM_199414 Homo sapiens protein regulator of cytokinesis 1 (PRC1), transcript variant 3,
 NM_199415 Homo sapiens likely ortholog of mouse ubiquitin conjugating enzyme 7 inte
 NM_199416 Homo sapiens papillary renal cell carcinoma (translocation-associated) (PRC
 NM_199417 Homo sapiens nuclear protein E3-3 (DKFZP564J0123), transcript variant 5, i
 NM_199418 Homo sapiens prolylcarboxypeptidase (angiotensinase C) (PRCP), transcript
 NM_199420 Homo sapiens polymerase (DNA directed), theta (POLQ), transcript variant 2
 NM_199421 Homo sapiens suppressor of cytokine signaling 4 (SOCS4), transcript variant
 NM_199423 Homo sapiens WW domain containing E3 ubiquitin protein ligase 2 (WWP2)
 NM_199424 Homo sapiens WW domain containing E3 ubiquitin protein ligase 2 (WWP2)
 NM_199425 Homo sapiens visual system homeobox 1 homolog, CHX10-like (zebrafish) (C
 NM_199426 Homo sapiens zinc finger protein 64 homolog (mouse) (ZFP64), transcript va
 NM_199427 Homo sapiens zinc finger protein 64 homolog (mouse) (ZFP64), transcript va
 NM_199436 Homo sapiens spastic paraplegia 4 (autosomal dominant; spastin) (SPG4), t
 NM_199437 Homo sapiens PR domain containing 10 (PRDM10), transcript variant 2, mR
 NM_199438 Homo sapiens PR domain containing 10 (PRDM10), transcript variant 3, mR
 NM_199439 Homo sapiens PR domain containing 10 (PRDM10), transcript variant 4, mR
 NM_199440 Homo sapiens heat shock 60kDa protein 1 (chaperonin) (HSPD1), nuclear g
 NM_199441 Homo sapiens zinc finger protein 334 (ZNF334), transcript variant 2, mRNA
 NM_199442 Homo sapiens coatomer protein complex, subunit epsilon (COPE), transcript
 NM_199443 Homo sapiens ubiquitin specific protease 4 (proto-oncogene) (USP4), transc

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NM_199444 Homo sapiens coatomer protein complex, subunit epsilon (COPE), transcript
 NM_199450 Homo sapiens zinc finger protein 365 (ZNF365), transcript variant B, mRNA
 NM_199451 Homo sapiens zinc finger protein 365 (ZNF365), transcript variant C, mRNA
 NM_199452 Homo sapiens zinc finger protein 365 (ZNF365), transcript variant D, mRNA
 NM_199453 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 4 (HTR4), transcript
 NM_199454 Homo sapiens PR domain containing 16 (PRDM16), transcript variant 2, mRNA
 NM_199456 Homo sapiens testis/prostate/placenta-expressed protein (TEPP), transcript
 NM_199459 Homo sapiens chromosome 10 open reading frame 71 (C10orf71), mRNA
 NM_199460 Homo sapiens hypothetical protein LOC283439 (LOC283439), mRNA
 NM_199461 Homo sapiens nanos homolog 1 (Drosophila) (NANOS1), mRNA
 NM_199462 Homo sapiens receptor interacting protein kinase 5 (RIPK5), transcript variant
 NM_199464 Homo sapiens potassium channel regulator (KCNRG), mRNA
 NM_199478 Homo sapiens proteolipid protein 1 (Pelizaeus-Merzbacher disease, spastic)
 NM_199482 Homo sapiens preimplantation protein 3 (PREI3), transcript variant 2, mRNA
 NM_199483 Homo sapiens chromosome 20 open reading frame 24 (C20orf24), transcript
 NM_199484 Homo sapiens chromosome 20 open reading frame 24 (C20orf24), transcript
 NM_199485 Homo sapiens chromosome 20 open reading frame 24 (C20orf24), transcript
 NM_199487 Homo sapiens chromosome 20 open reading frame 44 (C20orf44), transcript
 NM_199511 Homo sapiens steroid sensitive gene 1 (URB), transcript variant 1, mRNA
 NM_199512 Homo sapiens steroid sensitive gene 1 (URB), transcript variant 2, mRNA
 NM_199513 Homo sapiens chromosome 20 open reading frame 44 (C20orf44), transcript
 NM_201222 Homo sapiens melanoma antigen, family D, 2 (MAGED2), transcript variant 2
 NM_201224 Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 47 (DDX47), transcript
 NM_201252 Homo sapiens aflatoxin B1 aldehyde reductase 3 (AFAR3), mRNA
 NM_201253 Homo sapiens crumbs homolog 1 (Drosophila) (CRB1), transcript variant 2, mRNA
 NM_201259 Homo sapiens homolog of yeast TIM14 (TIM14), transcript variant 2, mRNA
 NM_201260 Homo sapiens homolog of yeast TIM14 (TIM14), transcript variant 3, mRNA
 NM_201261 Homo sapiens homolog of yeast TIM14 (TIM14), transcript variant 4, mRNA
 NM_201262 Homo sapiens DnaJ (Hsp40) homolog, subfamily C, member 12 (DNAJC12), mRNA
 NM_201263 Homo sapiens tryptophanyl tRNA synthetase 2 (mitochondrial) (WARS2), mRNA
 NM_201264 Homo sapiens neuropilin 2 (NRP2), transcript variant 6, mRNA
 NM_201265 Homo sapiens bone marrow stromal cell-derived ubiquitin-like (BMSCL-UBP),
 NM_201266 Homo sapiens neuropilin 2 (NRP2), transcript variant 1, mRNA
 NM_201267 Homo sapiens neuropilin 2 (NRP2), transcript variant 5, mRNA
 NM_201268 Homo sapiens membrane-bound transcription factor protease, site 1 (MBTPS1)
 NM_201269 Homo sapiens zinc finger motif enhancer binding protein 2 (Zep-2), transcript
 NM_201274 Homo sapiens myosin phosphatase-Rho interacting protein (M-RIP), mRNA
 NM_201277 Homo sapiens calponin 2 (CNN2), transcript variant 2, mRNA
 NM_201278 Homo sapiens myotubularin related protein 2 (MTMR2), transcript variant 2, mRNA
 NM_201279 Homo sapiens neuropilin 2 (NRP2), transcript variant 3, mRNA
 NM_201280 Homo sapiens muted homolog (mouse) (MUTED), mRNA
 NM_201281 Homo sapiens myotubularin related protein 2 (MTMR2), transcript variant 3, mRNA
 NM_201282 Homo sapiens epidermal growth factor receptor (erythroblastic leukemia virus)
 NM_201283 Homo sapiens epidermal growth factor receptor (erythroblastic leukemia virus)
 NM_201284 Homo sapiens epidermal growth factor receptor (erythroblastic leukemia virus)
 NM_201286 Homo sapiens ubiquitin specific protease 51 (USP51), mRNA
 NM_201348 Homo sapiens proline arginine-rich end leucine-rich repeat protein (PRELP),
 NM_201349 Homo sapiens docking protein 2, 56kDa (DOK2), transcript variant 2, mRNA
 NM_201377 Homo sapiens cancer susceptibility candidate 2 (CASC2), mRNA
 NM_201378 Homo sapiens plectin 1, intermediate filament binding protein 500kDa (PLEC
 NM_201379 Homo sapiens plectin 1, intermediate filament binding protein 500kDa (PLEC
 NM_201380 Homo sapiens plectin 1, intermediate filament binding protein 500kDa (PLEC
 NM_201381 Homo sapiens plectin 1, intermediate filament binding protein 500kDa (PLEC
 NM_201382 Homo sapiens plectin 1, intermediate filament binding protein 500kDa (PLEC
 NM_201383 Homo sapiens plectin 1, intermediate filament binding protein 500kDa (PLEC
 NM_201384 Homo sapiens plectin 1, intermediate filament binding protein 500kDa (PLEC
 NM_201397 Homo sapiens glutathione peroxidase 1 (GPX1), transcript variant 2, mRNA

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NM_201398 Homo sapiens FAD-synthetase (PP591), transcript variant 2, mRNA
 NM_201399 Homo sapiens DPH2-like 2 (S. cerevisiae) (DPH2L2), transcript variant 2, m
 NM_201400 Homo sapiens hypothetical protein SB153 (SB153), transcript variant 1, mR
 NM_201402 Homo sapiens deubiquitinating enzyme 3 (DUB3), mRNA
 NM_201403 Homo sapiens MOB1, Mps One Binder kinase activator-like 2C (yeast) (MOE
 NM_201412 Homo sapiens LUC7-like (S. cerevisiae) (LUC7L), transcript variant 2, mRNA
 NM_201413 Homo sapiens amyloid beta (A4) precursor protein (protease nexin-II, Alzhei
 NM_201414 Homo sapiens amyloid beta (A4) precursor protein (protease nexin-II, Alzhei
 NM_201428 Homo sapiens reticulon 3 (RTN3), transcript variant 2, mRNA
 NM_201429 Homo sapiens reticulon 3 (RTN3), transcript variant 3, mRNA
 NM_201430 Homo sapiens reticulon 3 (RTN3), transcript variant 4, mRNA
 NM_201431 Homo sapiens Ras association (RalGDS/AF-6) domain family 6 (RASSF6), t
 NM_201432 Homo sapiens growth arrest-specific 7 (GAS7), transcript variant b, mRNA
 NM_201433 Homo sapiens growth arrest-specific 7 (GAS7), transcript variant c, mRNA
 NM_201434 Homo sapiens RAB5C, member RAS oncogene family (RAB5C), transcript v
 NM_201435 Homo sapiens testis-specific protein TSP-NY (TSP-NY), transcript variant 2,
 NM_201436 Homo sapiens H2A histone family, member V (H2AFV), transcript variant 3, i
 NM_201437 Homo sapiens transcription elongation factor A (SII), 1 (TCEA1), transcript v
 NM_201438 Homo sapiens periphilin 1 (PPHLN1), transcript variant 5, mRNA
 NM_201439 Homo sapiens periphilin 1 (PPHLN1), transcript variant 3, mRNA
 NM_201440 Homo sapiens periphilin 1 (PPHLN1), transcript variant 4, mRNA
 NM_201441 Homo sapiens TEA domain family member 4 (TEAD4), transcript variant 2, n
 NM_201442 Homo sapiens complement component 1, s subcomponent (C1S), transcript
 NM_201443 Homo sapiens TEA domain family member 4 (TEAD4), transcript variant 3, n
 NM_201444 Homo sapiens diacylglycerol kinase, alpha 80kDa (DGKA), transcript variant
 NM_201445 Homo sapiens diacylglycerol kinase, alpha 80kDa (DGKA), transcript variant
 NM_201446 Homo sapiens EGF-like-domain, multiple 7 (EGFL7), transcript variant 2, mR
 NM_201453 Homo sapiens dopamine responsive protein (LOC220869), mRNA
 NM_201515 Homo sapiens periphilin 1 (PPHLN1), transcript variant 2, mRNA
 NM_201516 Homo sapiens H2A histone family, member V (H2AFV), transcript variant 4, i
 NM_201517 Homo sapiens H2A histone family, member V (H2AFV), transcript variant 5, i
 NM_201520 Homo sapiens similar to 1810012H11 Rik (FLJ40217), mRNA
 NM_201521 Homo sapiens kinesin-like 8 (KNSL8), transcript variant 1, mRNA
 NM_201522 Homo sapiens kinesin-like 8 (KNSL8), transcript variant 2, mRNA
 NM_201523 Homo sapiens kinesin-like 8 (KNSL8), transcript variant 3, mRNA
 NM_201524 Homo sapiens G protein-coupled receptor 56 (GPR56), transcript variant 2, r
 NM_201525 Homo sapiens G protein-coupled receptor 56 (GPR56), transcript variant 3, r
 NM_201526 Homo sapiens immunoglobulin superfamily containing leucine-rich repeat (IS
 NM_201532 Homo sapiens diacylglycerol kinase, zeta 104kDa (DGKZ), transcript variant
 NM_201533 Homo sapiens diacylglycerol kinase, zeta 104kDa (DGKZ), transcript variant
 NM_201535 Homo sapiens NDRG family member 2 (NDRG2), transcript variant 1, mRNA
 NM_201536 Homo sapiens NDRG family member 2 (NDRG2), transcript variant 3, mRNA
 NM_201537 Homo sapiens NDRG family member 2 (NDRG2), transcript variant 4, mRNA
 NM_201538 Homo sapiens NDRG family member 2 (NDRG2), transcript variant 5, mRNA
 NM_201539 Homo sapiens NDRG family member 2 (NDRG2), transcript variant 6, mRNA
 NM_201540 Homo sapiens NDRG family member 2 (NDRG2), transcript variant 7, mRNA
 NM_201541 Homo sapiens NDRG family member 2 (NDRG2), transcript variant 8, mRNA
 NM_201542 Homo sapiens mediator of RNA polymerase II transcription, subunit 8 homol
 NM_201543 Homo sapiens lectin, galactoside-binding, soluble, 8 (galectin 8) (LGALS8), t
 NM_201544 Homo sapiens lectin, galactoside-binding, soluble, 8 (galectin 8) (LGALS8), t
 NM_201545 Homo sapiens lectin, galactoside-binding, soluble, 8 (galectin 8) (LGALS8), t
 NM_201546 Homo sapiens hypothetical protein LOC200008 (LOC200008), mRNA
 NM_201548 Homo sapiens retinitis pigmentosa 26 (autosomal recessive) (RP26), mRNA
 NM_201550 Homo sapiens leucine rich repeat containing 10 (LRRC10), mRNA
 NM_201552 Homo sapiens fibrinogen-like 1 (FGL1), transcript variant 3, mRNA
 NM_201553 Homo sapiens fibrinogen-like 1 (FGL1), transcript variant 4, mRNA
 NM_201554 Homo sapiens diacylglycerol kinase, alpha 80kDa (DGKA), transcript variant

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NM_201555 Homo sapiens four and a half LIM domains 2 (FHL2), transcript variant 2, mRNA
 NM_201556 Homo sapiens four and a half LIM domains 2 (FHL2), transcript variant 3, mRNA
 NM_201557 Homo sapiens four and a half LIM domains 2 (FHL2), transcript variant 4, mRNA
 NM_201559 Homo sapiens forkhead box O3A (FOXO3A), transcript variant 2, mRNA
 NM_201563 Homo sapiens Fc fragment of IgG, low affinity IIc, receptor for (CD32) (FCGF)
 NM_201564 Homo sapiens chromosome 10 open reading frame 94 (C10orf94), mRNA
 NM_201565 Homo sapiens hypothetical gene supported by BC039313 (LOC284861), mRNA
 NM_201566 Homo sapiens solute carrier family 16 (monocarboxylic acid transporters), mRNA
 NM_201567 Homo sapiens cell division cycle 25A (CDC25A), transcript variant 2, mRNA
 NM_201568 Homo sapiens chromosome 1 open reading frame 16 (C1orf16), transcript variant 1
 NM_201569 Homo sapiens chromosome 1 open reading frame 16 (C1orf16), transcript variant 2
 NM_201570 Homo sapiens calcium channel, voltage-dependent, beta 2 subunit (CACNB2)
 NM_201571 Homo sapiens calcium channel, voltage-dependent, beta 2 subunit (CACNB2)
 NM_201572 Homo sapiens calcium channel, voltage-dependent, beta 2 subunit (CACNB2)
 NM_201574 Homo sapiens solute carrier family 4, anion exchanger, member 3 (SLC4A3)
 NM_201575 Homo sapiens seizure related 6 homolog (mouse)-like 2 (SEZ6L2), mRNA
 NM_201589 Homo sapiens v-maf musculoaponeurotic fibrosarcoma oncogene homolog 1
 NM_201590 Homo sapiens calcium channel, voltage-dependent, beta 2 subunit (CACNB2)
 NM_201591 Homo sapiens glycoprotein M6A (GPM6A), transcript variant 2, mRNA
 NM_201592 Homo sapiens glycoprotein M6A (GPM6A), transcript variant 3, mRNA
 NM_201593 Homo sapiens calcium channel, voltage-dependent, beta 2 subunit (CACNB2)
 NM_201594 Homo sapiens similar to B-cell linker; B cell linker protein (LOC284948), transcript
 NM_201595 Homo sapiens general transcription factor IIA, 1, 19/37kDa (GTF2A1), transcript
 NM_201596 Homo sapiens calcium channel, voltage-dependent, beta 2 subunit (CACNB2)
 NM_201597 Homo sapiens calcium channel, voltage-dependent, beta 2 subunit (CACNB2)
 NM_201598 Homo sapiens hypothetical protein SB153 (SB153), transcript variant 2, mRNA
 NM_201599 Homo sapiens zinc finger protein 261 (ZNF261), transcript variant 2, mRNA
 NM_201612 Homo sapiens IKK interacting protein (IKIP), transcript variant 2, mRNA
 NM_201613 Homo sapiens IKK interacting protein (IKIP), transcript variant 3.1, mRNA
 NM_201614 Homo sapiens IKK interacting protein (IKIP), transcript variant 3.2, mRNA
 NM_201623 Homo sapiens myeloid inhibitory C-type lectin-like receptor (MILCL), transcript
 NM_201624 Homo sapiens ubiquitin specific protease 33 (USP33), transcript variant 2, mRNA
 NM_201625 Homo sapiens myeloid inhibitory C-type lectin-like receptor (MILCL), transcript
 NM_201626 Homo sapiens ubiquitin specific protease 33 (USP33), transcript variant 3, mRNA
 NM_201627 Homo sapiens tripartite motif-containing 41 (TRIM41), transcript variant 2, mRNA
 NM_201628 Homo sapiens hypothetical protein FLJ43806 (FLJ43806), mRNA
 NM_201629 Homo sapiens tight junction protein 2 (zona occludens 2) (TJP2), transcript variant
 NM_201630 Homo sapiens leucine rich repeat neuronal 5 (LRRN5), transcript variant 2, mRNA
 NM_201631 Homo sapiens transglutaminase 5 (TGM5), transcript variant 1, mRNA
 NM_201632 Homo sapiens transcription factor 7 (T-cell specific, HMG-box) (TCF7), transcript
 NM_201633 Homo sapiens transcription factor 7 (T-cell specific, HMG-box) (TCF7), transcript
 NM_201634 Homo sapiens transcription factor 7 (T-cell specific, HMG-box) (TCF7), transcript
 NM_201636 Homo sapiens thromboxane A2 receptor (TBXA2R), transcript variant 1, mRNA
 NM_201647 Homo sapiens STAM binding protein (STAMPB), transcript variant 2, mRNA
 NM_201648 Homo sapiens glycine-N-acyltransferase (GLYAT), nuclear gene encoding mRNA
 NM_201649 Homo sapiens solute carrier family 6 (neurotransmitter transporter, glycine), mRNA
 NM_201650 Homo sapiens B7 gene (B7), transcript variant 1, mRNA
 NM_201651 Homo sapiens solute carrier family 28 (sodium-coupled nucleoside transporter)
 NM_201653 Homo sapiens eosinophil chemotactic cytokine (CHIA), mRNA
 NM_201694 Homo sapiens vesicle-associated membrane protein 4 (VAMP4), transcript variant
 NM_201695 Homo sapiens splicing factor 1 (SF1), transcript variant 2, mRNA
 NM_201697 Homo sapiens splicing factor 1 (SF1), transcript variant 4, mRNA
 NM_201698 Homo sapiens splicing factor 1 (SF1), transcript variant 3, mRNA
 NM_201699 Homo sapiens E74-like factor 2 (ets domain transcription factor) (ELF2), transcript
 NM_202000 Homo sapiens SA hypertension-associated homolog (rat) (SAH), transcript variant
 NM_202001 Homo sapiens excision repair cross-complementing rodent repair deficiency,
 NM_202002 Homo sapiens forkhead box M1 (FOXM1), transcript variant 1, mRNA

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NM_202003 Homo sapiens forkhead box M1 (FOXM1), transcript variant 3, mRNA
 NM_202004 Homo sapiens hemochromatosis type 2 (juvenile) (HFE2), transcript variant c
 NM_202467 Homo sapiens regulator of G-protein signalling 19 interacting protein 1 (RGS
 NM_202468 Homo sapiens regulator of G-protein signalling 19 interacting protein 1 (RGS
 NM_202469 Homo sapiens regulator of G-protein signalling 19 interacting protein 1 (RGS
 NM_202470 Homo sapiens regulator of G-protein signalling 19 interacting protein 1 (RGS
 NM_202494 Homo sapiens regulator of G-protein signalling 19 interacting protein 1 (RGS
 NM_202768 Homo sapiens rTS beta protein (HSRTSBETA), transcript variant 1, mRNA
 NM_203281 Homo sapiens BMX non-receptor tyrosine kinase (BMX), mRNA
 NM_203282 Homo sapiens zinc finger protein 539 (ZNF539), mRNA
 NM_203283 Homo sapiens recombining binding protein suppressor of hairless (Drosophil
 NM_203284 Homo sapiens recombining binding protein suppressor of hairless (Drosophil
 NM_203285 Homo sapiens poliovirus receptor-related 1 (herpesvirus entry mediator C; ne
 NM_203286 Homo sapiens poliovirus receptor-related 1 (herpesvirus entry mediator C; ne
 NM_203287 Homo sapiens pregnancy specific beta-1-glycoprotein 11 (PSG11), transcript
 NM_203288 Homo sapiens retinitis pigmentosa 9 (autosomal dominant) (RP9), mRNA
 NM_203289 Homo sapiens POU domain, class 5, transcription factor 1 (POU5F1), transc
 NM_203290 Homo sapiens polymerase (RNA) I polypeptide C, 30kDa (POLR1C), transcr
 NM_203291 Homo sapiens retinoblastoma binding protein 8 (RBBP8), transcript variant 2
 NM_203292 Homo sapiens retinoblastoma binding protein 8 (RBBP8), transcript variant 3
 NM_203293 Homo sapiens tripartite motif-containing 7 (TRIM7), transcript variant 1, mRN
 NM_203294 Homo sapiens tripartite motif-containing 7 (TRIM7), transcript variant 5, mRN
 NM_203295 Homo sapiens tripartite motif-containing 7 (TRIM7), transcript variant 4, mRN
 NM_203296 Homo sapiens tripartite motif-containing 7 (TRIM7), transcript variant 3, mRN
 NM_203297 Homo sapiens tripartite motif-containing 7 (TRIM7), transcript variant 2, mRN
 NM_203298 Homo sapiens coiled-coil-helix-coiled-coil-helix domain containing 1 (CHCHL
 NM_203299 Homo sapiens hypothetical protein MGC41945 (MGC41945), mRNA
 NM_203301 Homo sapiens F-box protein 33 (FBXO33), mRNA
 NM_203302 Homo sapiens similar to RPL23AP7 protein (MGC70863), transcript variant 2
 NM_203303 Homo sapiens cellular nucleic acid binding protein-like (LOC389874), mRNA
 NM_203304 Homo sapiens ring finger and KH domain containing 1 (RKH1), mRNA
 NM_203305 Homo sapiens hypothetical protein MGC50853 (MGC50853), mRNA
 NM_203306 Homo sapiens hypothetical protein MGC39606 (MGC39606), mRNA
 NM_203307 Homo sapiens hypothetical protein MGC35402 (MGC35402), mRNA
 NM_203308 Homo sapiens ribosomal protein L13A-like (MGC34774), mRNA
 NM_203309 Homo sapiens hypothetical MGC48595 (MGC48595), mRNA
 NM_203311 Homo sapiens similar to Taxol resistant associated protein 3 (TRAG-3) (LOC
 NM_203314 Homo sapiens 3-hydroxybutyrate dehydrogenase (heart, mitochondrial) (BDH
 NM_203315 Homo sapiens 3-hydroxybutyrate dehydrogenase (heart, mitochondrial) (BDH
 NM_203316 Homo sapiens dolichyl-phosphate (UDP-N-acetylglucosamine) N-acetylgluco
 NM_203318 Homo sapiens myosin XVIIIa (MYO18A), transcript variant 2, mRNA
 NM_203326 Homo sapiens 5-azacytidine induced 2 (AZI2), transcript variant 2, mRNA
 NM_203327 Homo sapiens solute carrier family 23 (nucleobase transporters), member 2
 NM_203329 Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antib
 NM_203330 Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antib
 NM_203331 Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antib
 NM_203339 Homo sapiens clusterin (complement lysis inhibitor, SP-40,40, sulfated glyco
 NM_203341 Homo sapiens 15 kDa selenoprotein (SEP15), transcript variant 2, mRNA
 NM_203342 Homo sapiens erythrocyte membrane protein band 4.1 (elliptocytosis 1, RH-I
 NM_203343 Homo sapiens erythrocyte membrane protein band 4.1 (elliptocytosis 1, RH-I
 NM_203344 Homo sapiens SERTA domain containing 3 (SERTA3), transcript variant 2,
 NM_203346 Homo sapiens high density lipoprotein binding protein (vlgilin) (HDLBP), mR
 NM_203347 Homo sapiens MSFL2541 (UNQ2541), mRNA
 NM_203348 Homo sapiens hypothetical MGC50722 (MGC50722), mRNA
 NM_203349 Homo sapiens ral-like protein (RalP), mRNA
 NM_203350 Homo sapiens zinc finger protein 265 (ZNF265), transcript variant 1, mRNA
 NM_203351 Homo sapiens mitogen-activated protein kinase kinase kinase 3 (MAP3K3), l

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NM_203352 Homo sapiens PDZ and LIM domain 7 (enigma) (PDLIM7), transcript variant
 NM_203353 Homo sapiens PDZ and LIM domain 7 (enigma) (PDLIM7), transcript variant
 NM_203354 Homo sapiens meningioma expressed antigen 6 (coiled-coil proline-rich) (MC
 NM_203355 Homo sapiens meningioma expressed antigen 6 (coiled-coil proline-rich) (MC
 NM_203356 Homo sapiens meningioma expressed antigen 6 (coiled-coil proline-rich) (MC
 NM_203357 Homo sapiens meningioma expressed antigen 6 (coiled-coil proline-rich) (MC
 NM_203364 Homo sapiens membrane component, chromosome 11, surface marker 1 (M
 NM_203365 Homo sapiens Ras association (RalGDS/AF-6) and pleckstrin homology dom
 NM_203370 Homo sapiens similar to RIKEN cDNA 6530416L21 (LOC389119), mRNA
 NM_203371 Homo sapiens similar to RIKEN cDNA 1110018M03 (LOC387758), mRNA
 NM_203372 Homo sapiens acyl-CoA synthetase long-chain family member 3 (ACSL3), tr
 NM_203373 Homo sapiens hypothetical protein LOC283807 (LOC283807), mRNA
 NM_203374 Homo sapiens similar to zinc finger protein (LOC147808), mRNA
 NM_203375 Homo sapiens hypothetical gene supported by BC001801 (LOC284912), mR
 NM_203376 Homo sapiens similar to RIKEN cDNA 4930429O20 (LOC388730), mRNA
 NM_203377 Homo sapiens myoglobin (MB), transcript variant 2, mRNA
 NM_203378 Homo sapiens myoglobin (MB), transcript variant 3, mRNA
 NM_203379 Homo sapiens acyl-CoA synthetase long-chain family member 5 (ACSL5), tr
 NM_203380 Homo sapiens acyl-CoA synthetase long-chain family member 5 (ACSL5), tr
 NM_203381 Homo sapiens protein for MGC71805 (MGC71805), mRNA
 NM_203382 Homo sapiens alpha-methylacyl-CoA racemase (AMACR), transcript variant
 NM_203383 Homo sapiens ribonuclease/angiogenin inhibitor (RNH), transcript variant 2, i
 NM_203384 Homo sapiens ribonuclease/angiogenin inhibitor (RNH), transcript variant 3, i
 NM_203385 Homo sapiens ribonuclease/angiogenin inhibitor (RNH), transcript variant 4, i
 NM_203386 Homo sapiens ribonuclease/angiogenin inhibitor (RNH), transcript variant 5, i
 NM_203387 Homo sapiens ribonuclease/angiogenin inhibitor (RNH), transcript variant 6, i
 NM_203388 Homo sapiens ribonuclease/angiogenin inhibitor (RNH), transcript variant 7, i
 NM_203389 Homo sapiens ribonuclease/angiogenin inhibitor (RNH), transcript variant 8, i
 NM_203390 Homo sapiens similar to RIKEN cDNA 3000004N20 (LOC389677), mRNA
 NM_203391 Homo sapiens glycerol kinase (GK), transcript variant 1, mRNA
 NM_203392 Homo sapiens hypothetical gene supported by BC031617 (LOC284123), mR
 NM_203393 Homo sapiens hypothetical gene supported by BC031661 (LOC389458), mR
 NM_203394 Homo sapiens E2F transcription factor 7 (E2F7), mRNA
 NM_203395 Homo sapiens iodotyrosine dehalogenase 1 (DEHAL1), mRNA
 NM_203399 Homo sapiens stathmin 1/oncoprotein 18 (STMN1), transcript variant 2, mR
 NM_203400 Homo sapiens similar to candidate mediator of the p53-dependent G2 arrest
 NM_203401 Homo sapiens stathmin 1/oncoprotein 18 (STMN1), transcript variant 1, mR
 NM_203402 Homo sapiens similar to CG10671-like (LOC161247), mRNA
 NM_203403 Homo sapiens chromosome 9 open reading frame 150 (C9orf150), mRNA
 NM_203405 Homo sapiens similar to RIKEN cDNA 2310002B14 (LOC388818), mRNA
 NM_203406 Homo sapiens similar to metallo-beta-lactamase superfamily protein (LOC15
 NM_203407 Homo sapiens similar to CG32656-PA (LOC340602), mRNA
 NM_203408 Homo sapiens similar to hypothetical protein FLJ35782 (LOC158724), mRN/
 NM_203411 Homo sapiens similar to RIKEN cDNA 2600017H02 (LOC92162), mRNA
 NM_203412 Homo sapiens similar to RIKEN cDNA 4930522D07 (LOC164153), mRNA
 NM_203413 Homo sapiens S-phase 2 protein (DERP6), mRNA
 NM_203414 Homo sapiens S-phase 2 protein (DERP6), mRNA
 NM_203415 Homo sapiens S-phase 2 protein (DERP6), mRNA
 NM_203416 Homo sapiens CD163 antigen (CD163), transcript variant 2, mRNA
 NM_203417 Homo sapiens Down syndrome critical region gene 1 (DSCR1), transcript vai
 NM_203418 Homo sapiens Down syndrome critical region gene 1 (DSCR1), transcript vai
 NM_203419 Homo sapiens hypothetical protein LOC286016 (LOC286016), mRNA
 NM_203422 Homo sapiens similar to hypothetical protein (LOC221091), mRNA
 NM_203423 Homo sapiens hypothetical gene supported by BC031673 (LOC389199), mR
 NM_203424 Homo sapiens hypothetical protein MGC50809 (LOC389123), mRNA
 NM_203425 Homo sapiens hypothetical gene supported by BC046200 (LOC388407), mR
 NM_203426 Homo sapiens hypothetical protein FLJ0297 (LOC388152), mRNA

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NM_203428 Homo sapiens Down syndrome critical region gene 8 (DSCR8), transcript vai
 NM_203429 Homo sapiens Down syndrome critical region gene 8 (DSCR8), transcript vai
 NM_203430 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), transcript va
 NM_203431 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), transcript va
 NM_203433 Homo sapiens Down syndrome critical region gene 2 (DSCR2), transcript vai
 NM_203434 Homo sapiens similar to RIKEN cDNA 2610524G09 (LOC389792), mRNA
 NM_203436 Homo sapiens achaete-scute complex-like 4 (Drosophila) (ASCL4), mRNA
 NM_203437 Homo sapiens atrophilin protein (AFTIPHILIN), transcript variant 1, mRNA
 NM_203438 Homo sapiens chromosome 10 open reading frame 4 (C10orf4), transcript va
 NM_203439 Homo sapiens chromosome 10 open reading frame 4 (C10orf4), transcript va
 NM_203440 Homo sapiens chromosome 10 open reading frame 4 (C10orf4), transcript va
 NM_203441 Homo sapiens chromosome 10 open reading frame 4 (C10orf4), transcript va
 NM_203444 Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 9 (A
 NM_203445 Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 9 (A
 NM_203446 Homo sapiens synaptotagmin 1 (SYNJ1), transcript variant 2, mRNA
 NM_203447 Homo sapiens dedicator of cytokinesis 8 (DOCK8), mRNA
 NM_203448 Homo sapiens hypothetical protein MGC21881 (MGC21881), mRNA
 NM_203451 Homo sapiens hypothetical LOC400120 (LOC400120), mRNA
 NM_203452 Homo sapiens hypothetical protein LOC403312 (MGC39545), mRNA
 NM_203453 Homo sapiens hypothetical LOC403313 (LOC403313), mRNA
 NM_203454 Homo sapiens hypothetical protein MGC26594 (MGC26594), mRNA
 NM_203456 Homo sapiens peptidylprolyl isomerase E (cyclophilin E) (PPIE), transcript va
 NM_203457 Homo sapiens peptidylprolyl isomerase E (cyclophilin E) (PPIE), transcript va
 NM_203458 Homo sapiens similar to NOTCH2 protein (N2N), mRNA
 NM_203459 Homo sapiens KIAA1078 protein (KIAA1078), mRNA
 NM_203462 Homo sapiens PP784 protein (PP784), transcript variant 2, mRNA
 NM_203463 Homo sapiens LAG1 longevity assurance homolog 6 (S. cerevisiae) (LASS6)
 NM_203466 Homo sapiens peptidylprolyl isomerase (cyclophilin) like 5 (PPII5), transcript
 NM_203467 Homo sapiens peptidylprolyl isomerase (cyclophilin) like 5 (PPII5), transcript
 NM_203468 Homo sapiens ectonucleoside triphosphate diphosphohydrolase 2 (ENTPD2,
 NM_203471 Homo sapiens lectin, galactoside-binding, soluble, 14 (LGALS14), transcript
 NM_203472 Homo sapiens selenoprotein S (SELS), transcript variant 1, mRNA
 NM_203473 Homo sapiens porcupine homolog (Drosophila) (PORCN), transcript variant I
 NM_203474 Homo sapiens porcupine homolog (Drosophila) (PORCN), transcript variant I
 NM_203475 Homo sapiens porcupine homolog (Drosophila) (PORCN), transcript variant I
 NM_203476 Homo sapiens porcupine homolog (Drosophila) (PORCN), transcript variant I
 NM_203477 Homo sapiens similar to RPL23AP7 protein (MGC70863), transcript variant 1
 NM_203481 Homo sapiens hypothetical LOC4033340 (MGC70870), mRNA
 NM_203486 Homo sapiens delta-like 3 (Drosophila) (DLL3), transcript variant 2, mRNA
 NM_203487 Homo sapiens protocadherin 9 (PCDH9), transcript variant 1, mRNA
 NM_203488 Homo sapiens acylphosphatase 1, erythrocyte (common) type (ACYP1), tran
 NM_203494 Homo sapiens ubiquitin specific protease 50 (USP50), mRNA
 NM_203495 Homo sapiens COMM domain containing 6 (COMM6), transcript variant 2, i
 NM_203497 Homo sapiens COMM domain containing 6 (COMM6), transcript variant 1, i
 NM_203499 Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 42 (DDX42), transc
 NM_203500 Homo sapiens kelch-like ECH-associated protein 1 (KEAP1), transcript varia
 NM_203503 Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain
 NM_203504 Homo sapiens Ras-GTPase activating protein SH3 domain-binding protein 2
 NM_203505 Homo sapiens Ras-GTPase activating protein SH3 domain-binding protein 2
 NM_203506 Homo sapiens growth factor receptor-bound protein 2 (GRB2), transcript vari
 NM_203510 Homo sapiens transmembrane 6 superfamily member 2 (TM6SF2), transcript
 NM_205543 Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region,
 NM_205545 Homo sapiens RGTR430 (UNQ430), mRNA
 NM_205548 Homo sapiens AASA9217 (UNQ8217), mRNA
 NM_205767 Homo sapiens QIL1 protein (QIL1), mRNA
 NM_205768 Homo sapiens zinc finger protein 238 (ZNF238), transcript variant 1, mRNA
 NM_205833 Homo sapiens immunoglobulin superfamily, member 1 (IGSF1), transcript va

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NM_205834 Homo sapiens liver-specific bHLH-Zip transcription factor (LISCH7), transcript
 NM_205835 Homo sapiens liver-specific bHLH-Zip transcription factor (LISCH7), transcript
 NM_205836 Homo sapiens F-box protein 38 (FBXO38), transcript variant 2, mRNA
 NM_205837 Homo sapiens leukocyte specific transcript 1 (LST1), transcript variant 2, mR
 NM_205838 Homo sapiens leukocyte specific transcript 1 (LST1), transcript variant 3, mR
 NM_205839 Homo sapiens leukocyte specific transcript 1 (LST1), transcript variant 4, mR
 NM_205840 Homo sapiens leukocyte specific transcript 1 (LST1), transcript variant 5, mR
 NM_205841 Homo sapiens protease inhibitor H (MGC21394), mRNA
 NM_205842 Homo sapiens NCK-associated protein 1 (NCKAP1), transcript variant 2, mR
 NM_205843 Homo sapiens nuclear factor 1/C (CCAAT-binding transcription factor) (NFIC)
 NM_205845 Homo sapiens aldo-keto reductase family 1, member C2 (dihydrodiol dehydro
 NM_205846 Homo sapiens hypothetical protein MGC21644 (MGC21644), transcript varia
 NM_205847 Homo sapiens GDP-mannose pyrophosphorylase A (GMPPA), transcript vari
 NM_205848 Homo sapiens synaptotagmin VI (SYT6), mRNA
 NM_205849 Homo sapiens family with sequence similarity 9, member B (FAM9B), mRNA
 NM_205850 Homo sapiens solute carrier family 24, member 5 (SLC24A5), mRNA
 NM_205852 Homo sapiens macrophage antigen h (UNQ5782), mRNA
 NM_205853 Homo sapiens musculoskeletal, embryonic nuclear protein 1 (MUSTN1), mR
 NM_205854 Homo sapiens GSGL541 (UNQ541), mRNA
 NM_205855 Homo sapiens HWKM1940 (UNQ1940), mRNA
 NM_205856 Homo sapiens PNPK8288 (LOC389852), mRNA
 NM_205857 Homo sapiens FBI4 protein (FBI4), mRNA
 NM_205858 Homo sapiens neuromedin B (NMB), transcript variant 2, mRNA
 NM_205859 Homo sapiens olfactory receptor, family 2, subfamily K, member 2 (OR2K2),
 NM_205860 Homo sapiens nuclear receptor subfamily 5, group A, member 2 (NR5A2), tr
 NM_205861 Homo sapiens dehydrodolichyl diphosphate synthase (DHDDS), transcript va
 NM_205862 Homo sapiens UDP glycosyltransferase 1 family, polypeptide A6 (UGT1A6),
 NM_205863 Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region,
 NM_205864 Homo sapiens cancer/testis antigen 3 (CTAG3), transcript variant 2, mRNA
 NM_205865 Homo sapiens hypothetical protein LOC284381 (LOC284381), transcript vari
 NM_205869 Homo sapiens EGF-like-domain, multiple 9 (EGFL9), transcript variant 2, mR
 NM_205894 Homo sapiens estrogen-related receptor gamma (ESRRG), transcript vari
 NM_205895 Homo sapiens estrogen-related receptor gamma (ESRRG), transcript variant
 NM_208808 Homo sapiens citrate lyase beta like (CLYBL), transcript variant 2, mRNA
 NM_208809 Homo sapiens myelin oligodendrocyte glycoprotein (MOG), transcript variant
 NM_208810 Homo sapiens myelin oligodendrocyte glycoprotein (MOG), transcript variant
 NM_208811 Homo sapiens myelin oligodendrocyte glycoprotein (MOG), transcript variant
 NM_208812 Homo sapiens myelin oligodendrocyte glycoprotein (MOG), transcript variant
 NM_208813 Homo sapiens myelin oligodendrocyte glycoprotein (MOG), transcript variant
 NM_208814 Homo sapiens myelin oligodendrocyte glycoprotein (MOG), transcript variant
 NM_208817 Homo sapiens osteoclast-associated receptor (OSCAR), transcript variant 2,
 NM_208818 Homo sapiens osteoclast-associated receptor (OSCAR), transcript variant 1,
 NM_208819 Homo sapiens myosin binding protein C, slow type (MYBPC1), transcript vari
 NM_208820 Homo sapiens myosin binding protein C, slow type (MYBPC1), transcript vari
 NM_208821 Homo sapiens myosin binding protein C, slow type (MYBPC1), transcript vari
 NM_208824 Homo sapiens vitamin K epoxide reductase complex, subunit 1 (VKORC1), b
 NM_208825 Homo sapiens nucleostemin (NS), transcript variant 2, mRNA
 NM_208826 Homo sapiens nucleostemin (NS), transcript variant 3, mRNA
 NM_208827 Homo sapiens RAS-like, family 11, member A (RASL11A), mRNA
 NM_208828 Homo sapiens NACHT, leucine rich repeat and PYD containing 7 (NALP7), b
 NM_208831 Homo sapiens zinc finger, CSL domain containing 2 (ZCSSL2), mRNA
 NM_208832 Homo sapiens AWKS9372 (UNQ9372), mRNA
 NM_208833 Homo sapiens cortixin 1 (CTXN1), mRNA
 NM_208834 Homo sapiens chromosome 6 open reading frame 201 (C6orf201), mRNA
 NM_208835 Homo sapiens TNF receptor-associated factor 7 (TRAF7), transcript variant 2
 NM_208836 Homo sapiens peroxisomal D3,D2-enoyl-CoA isomerase (PECI), transcript v
 NM_208837 Homo sapiens oxidoredo-nitro domain-containing protein (NOR1), transcript v

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NM_206838 Homo sapiens similar to Hypothetical protein CBG21647 (LOC390511), mRNA
 NM_206839 Homo sapiens mortality factor 4 like 1 (MORF4L1), transcript variant 2, mRNA
 NM_206840 Homo sapiens nuclear VCP-like (NVL), transcript variant 2, mRNA
 NM_206841 Homo sapiens Fraser syndrome 1 (FRAS1), transcript variant 2, mRNA
 NM_206852 Homo sapiens reticulon 1 (RTN1), transcript variant 3, mRNA
 NM_206853 Homo sapiens quaking homolog, KH domain RNA binding (mouse) (QKI), tra
 NM_206854 Homo sapiens quaking homolog, KH domain RNA binding (mouse) (QKI), tra
 NM_206855 Homo sapiens quaking homolog, KH domain RNA binding (mouse) (QKI), tra
 NM_206857 Homo sapiens reticulon 1 (RTN1), transcript variant 2, mRNA
 NM_206858 Homo sapiens similar to protein phosphatase 1, regulatory (inhibitor) subunit
 NM_206860 Homo sapiens transforming, acidic coiled-coil containing protein 2 (TACC2),
 NM_206861 Homo sapiens transforming, acidic coiled-coil containing protein 2 (TACC2),
 NM_206862 Homo sapiens transforming, acidic coiled-coil containing protein 2 (TACC2),
 NM_206866 Homo sapiens BTB and CNC homology 1, basic leucine zipper transcription
 NM_206873 Homo sapiens protein phosphatase 1, catalytic subunit, alpha isoform (PPP1
 NM_206876 Homo sapiens protein phosphatase 1, catalytic subunit, beta isoform (PPP1C
 NM_206877 Homo sapiens protein phosphatase 1, catalytic subunit, beta isoform (PPP1C
 NM_206880 Homo sapiens olfactory receptor, family 2, subfamily V, member 2 (OR2V2),
 NM_206883 Homo sapiens prestin (motor protein) (PRES), transcript variant b, mRNA
 NM_206884 Homo sapiens prestin (motor protein) (PRES), transcript variant c, mRNA
 NM_206885 Homo sapiens prestin (motor protein) (PRES), transcript variant d, mRNA
 NM_206886 Homo sapiens sarcoma antigen NY-SAR-41 (NY-SAR-41), mRNA
 NM_206887 Homo sapiens Down syndrome cell adhesion molecule (DSCAM), transcript
 NM_206889 Homo sapiens chromosome 21 open reading frame 106 (C21orf106), transcr
 NM_206890 Homo sapiens chromosome 21 open reading frame 106 (C21orf106), transcr
 NM_206891 Homo sapiens chromosome 21 open reading frame 106 (C21orf106), transcr
 NM_206892 Homo sapiens malate dehydrogenase 1B, NAD (soluble) (MDH1B), mRNA
 NM_206893 Homo sapiens membrane-spanning 4-domains, subfamily A, member 10 (MS
 NM_206894 Homo sapiens hypothetical protein LOC388536 (MGC62100), mRNA
 NM_206895 Homo sapiens ASCL830 (UNQ830), mRNA
 NM_206898 Homo sapiens chromosome 21 open reading frame 61 (C21orf61), transcript
 NM_206900 Homo sapiens reticulon 2 (RTN2), transcript variant 2, mRNA
 NM_206901 Homo sapiens reticulon 2 (RTN2), transcript variant 3, mRNA
 NM_206902 Homo sapiens reticulon 2 (RTN2), transcript variant 4, mRNA
 NM_206907 Homo sapiens protein kinase, AMP-activated, alpha 1 catalytic subunit (PRK
 NM_206908 Homo sapiens chromosome 6 open reading frame 216 (C6orf216), transcript
 NM_206909 Homo sapiens pleckstrin and Sec7 domain containing 3 (PSD3), transcript vi
 NM_206910 Homo sapiens chromosome 6 open reading frame 216 (C6orf216), transcript
 NM_206911 Homo sapiens chromosome 6 open reading frame 216 (C6orf216), transcript
 NM_206912 Homo sapiens chromosome 6 open reading frame 216 (C6orf216), transcript
 NM_206914 Homo sapiens hepatocellular carcinoma-associated antigen HCA557a (DKF2
 NM_206915 Homo sapiens nerve growth factor receptor (TNFRSF16) associated protein
 NM_206917 Homo sapiens nerve growth factor receptor (TNFRSF16) associated protein
 NM_206918 Homo sapiens chromosome 14 open reading frame 66 (C14orf66), mRNA
 NM_206919 Homo sapiens ADP-ribosylation factor-like 9 (ARL9), mRNA
 NM_206920 Homo sapiens apical early endosomal glycoprotein precursor (AEGP), mRNA
 NM_206921 Homo sapiens chromosome 6 open reading frame 204 (C6orf204), mRNA
 NM_206922 Homo sapiens thymus LIM protein TLP-A (TLP), mRNA
 NM_206923 Homo sapiens YY2 transcription factor (YY2), mRNA
 NM_206925 Homo sapiens carbonic anhydrase XII (CA12), transcript variant 2, mRNA
 NM_206926 Homo sapiens selenoprotein N, 1 (SEPN1), transcript variant 2, mRNA
 NM_206927 Homo sapiens synaptotagmin-like 2 (SYTL2), transcript variant c, mRNA
 NM_206928 Homo sapiens synaptotagmin-like 2 (SYTL2), transcript variant d, mRNA
 NM_206929 Homo sapiens synaptotagmin-like 2 (SYTL2), transcript variant e, mRNA
 NM_206930 Homo sapiens synaptotagmin-like 2 (SYTL2), transcript variant f, mRNA
 NM_206933 Homo sapiens Usher syndrome 2A (autosomal recessive, mild) (USH2A), tra
 NM_206937 Homo sapiens ligase IV, DNA, ATP-dependent (LIG4), mRNA

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NM_206938 Homo sapiens membrane-spanning 4-domains, subfamily A, member 7 (MS-
 NM_206939 Homo sapiens membrane-spanning 4-domains, subfamily A, member 7 (MS-
 NM_206940 Homo sapiens membrane-spanning 4-domains, subfamily A, member 7 (MS-
 NM_206943 Homo sapiens latent transforming growth factor beta binding protein 1 (LTBF
 NM_206944 Homo sapiens transient receptor potential cation channel, subfamily M, mem
 NM_206945 Homo sapiens transient receptor potential cation channel, subfamily M, mem
 NM_206946 Homo sapiens transient receptor potential cation channel, subfamily M, mem
 NM_206947 Homo sapiens transient receptor potential cation channel, subfamily M, mem
 NM_206948 Homo sapiens transient receptor potential cation channel, subfamily M, mem
 NM_206949 Homo sapiens family with sequence similarity 14, member B (FAM14B), mRN
 NM_206953 Homo sapiens preferentially expressed antigen in melanoma (PRAE), trans
 NM_206954 Homo sapiens preferentially expressed antigen in melanoma (PRAE), trans
 NM_206955 Homo sapiens preferentially expressed antigen in melanoma (PRAE), trans
 NM_206956 Homo sapiens preferentially expressed antigen in melanoma (PRAE), trans
 NM_206961 Homo sapiens leukocyte tyrosine kinase (LTK), transcript variant 2, mRNA
 NM_206962 Homo sapiens HMT1 hnRNP methyltransferase-like 1 (S. cerevisiae) (HRMT
 NM_206963 Homo sapiens retinoic acid receptor responder (tazarotene Induced) 1 (RAR)
 NM_206964 Homo sapiens family with sequence similarity 3, member B (FAM3B), transcr
 NM_206965 Homo sapiens formiminotransferase cyclodeaminase (FTCD), transcript vari
 NM_206966 Homo sapiens similar to AVLV472 (MGC23985), mRNA
 NM_206967 Homo sapiens MGC17624 protein (MGC17624), mRNA
 NM_206994 Homo sapiens gonadotropin-releasing hormone (type 2) receptor 2 (GNRHR
 NM_206996 Homo sapiens projection protein PF6 (PF6), mRNA
 NM_206997 Homo sapiens G protein-coupled receptor 152 (GPR152), mRNA
 NM_206998 Homo sapiens secretoglobulin family 1D member 4 (SCGB1D4), mRNA
 NM_206999 Homo sapiens KIAA1007 protein (KIAA1007), transcript variant 2, mRNA
 NM_207002 Homo sapiens BCL2-like 11 (apoptosis facilitator) (BCL2L11), transcript vari
 NM_207003 Homo sapiens BCL2-like 11 (apoptosis facilitator) (BCL2L11), transcript vari
 NM_207005 Homo sapiens upstream transcription factor 1 (USF1), transcript variant 2, m
 NM_207006 Homo sapiens hypothetical protein MGC14128 (BJ-TSA-9), transcript variant
 NM_207007 Homo sapiens chemokine (C-C motif) ligand 4-like 1, telomeric (CCL4L1), m
 NM_207009 Homo sapiens family with sequence similarity 45, member A (FAM45A), mRN
 NM_207012 Homo sapiens adaptor-related protein complex 3, mu 1 subunit (AP3M1), tra
 NM_207013 Homo sapiens transcription elongation factor B (SII), polypeptide 2 (18kDa, i
 NM_207014 Homo sapiens hypothetical protein FLJ23129 (FLJ23129), transcript variant :
 NM_207015 Homo sapiens N-acetylated alpha-linked acidic dipeptidase 2 (NAALADL2), i
 NM_207032 Homo sapiens endothelin 3 (EDN3), transcript variant 2, mRNA
 NM_207033 Homo sapiens endothelin 3 (EDN3), transcript variant 3, mRNA
 NM_207034 Homo sapiens endothelin 3 (EDN3), transcript variant 4, mRNA
 NM_207035 Homo sapiens NP0014 protein (NP0014), transcript variant 1, mRNA
 NM_207036 Homo sapiens transcription factor 12 (HTF4, helix-loop-helix transcription fac
 NM_207037 Homo sapiens transcription factor 12 (HTF4, helix-loop-helix transcription fac
 NM_207038 Homo sapiens transcription factor 12 (HTF4, helix-loop-helix transcription fac
 NM_207040 Homo sapiens transcription factor 12 (HTF4, helix-loop-helix transcription fac
 NM_207042 Homo sapiens endosulfine alpha (ENSA), transcript variant 1, mRNA
 NM_207043 Homo sapiens endosulfine alpha (ENSA), transcript variant 2, mRNA
 NM_207044 Homo sapiens endosulfine alpha (ENSA), transcript variant 4, mRNA
 NM_207045 Homo sapiens endosulfine alpha (ENSA), transcript variant 5, mRNA
 NM_207046 Homo sapiens endosulfine alpha (ENSA), transcript variant 6, mRNA
 NM_207047 Homo sapiens endosulfine alpha (ENSA), transcript variant 7, mRNA
 NM_207102 Homo sapiens F-box and WD-40 domain protein 12 (FBXW12), mRNA
 NM_207103 Homo sapiens DTFT5783 (UNQ5783), mRNA
 NM_207106 Homo sapiens three prime repair exonuclease 2 (TREX2), transcript variant :
 NM_207107 Homo sapiens three prime repair exonuclease 2 (TREX2), transcript variant t
 NM_207108 Homo sapiens astrotactin (ASTN), transcript variant 2, mRNA
 NM_207111 Homo sapiens TRIAD3 protein (TRIAD3), transcript variant 1, mRNA
 NM_207112 Homo sapiens hydroxyacylglutathione hydrolase-like (HAGHL), transcript var

NM_207113 Homo sapiens solute carrier family 37 (glycerol-3-phosphate transporter), me
 NM_207115 Homo sapiens zinc finger protein 580 (ZNF580), transcript variant 2, mRNA
 NM_207116 Homo sapiens TRIAD3 protein (TRIAD3), transcript variant 2, mRNA
 NM_207117 Homo sapiens chromosome 14 open reading frame 68 (C14orf68), mRNA
 NM_207118 Homo sapiens general transcription factor IIH, polypeptide 5 (GTF2H5), mRNA
 NM_207119 Homo sapiens leucine rich repeat containing 20 (LRRC20), transcript variant
 NM_207121 Homo sapiens chromosome 20 open reading frame 55 (C20orf55), transcript
 NM_207122 Homo sapiens exostoses (multiple) 2 (EXT2), transcript variant 2, mRNA
 NM_207123 Homo sapiens GRB2-associated binding protein 1 (GAB1), transcript variant
 NM_207125 Homo sapiens polyamine oxidase (exo-N4-amino) (PAOX), transcript variant
 NM_207126 Homo sapiens polyamine oxidase (exo-N4-amino) (PAOX), transcript variant
 NM_207127 Homo sapiens polyamine oxidase (exo-N4-amino) (PAOX), transcript variant
 NM_207128 Homo sapiens polyamine oxidase (exo-N4-amino) (PAOX), transcript variant
 NM_207129 Homo sapiens polyamine oxidase (exo-N4-amino) (PAOX), transcript variant
 NM_207168 Homo sapiens endosulfine alpha (ENSA), transcript variant 8, mRNA
 NM_207170 Homo sapiens GCIP-interacting protein p29 (P29), transcript variant 2, mRNA
 NM_207171 Homo sapiens pogo transposable element with ZNF domain (POGZ), transcript
 NM_207172 Homo sapiens G protein-coupled receptor 154 (GPR154), transcript variant 1
 NM_207173 Homo sapiens G protein-coupled receptor 154 (GPR154), transcript variant 2
 NM_207174 Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 1 (ABC
 NM_207181 Homo sapiens nephronophthisis 1 (juvenile) (NPHP1), transcript variant 2, m
 NM_207186 Homo sapiens olfactory receptor, family 10, subfamily A, member 4 (OR10A
 NM_207189 Homo sapiens bromodomain, testis-specific (BRDT), transcript variant 1, mR
 NM_207191 Homo sapiens a disintegrin and metalloproteinase domain 15 (metargidin) (A
 NM_207194 Homo sapiens a disintegrin and metalloproteinase domain 15 (metargidin) (A
 NM_207195 Homo sapiens a disintegrin and metalloproteinase domain 15 (metargidin) (A
 NM_207196 Homo sapiens a disintegrin and metalloproteinase domain 15 (metargidin) (A
 NM_207197 Homo sapiens a disintegrin and metalloproteinase domain 15 (metargidin) (A
 NM_207283 Homo sapiens AAA1 protein (AAA1), transcript variant IX, mRNA
 NM_207284 Homo sapiens AAA1 protein (AAA1), transcript variant II, mRNA
 NM_207285 Homo sapiens AAA1 protein (AAA1), transcript variant III, mRNA
 NM_207286 Homo sapiens AAA1 protein (AAA1), transcript variant IV, mRNA
 NM_207287 Homo sapiens AAA1 protein (AAA1), transcript variant V, mRNA
 NM_207288 Homo sapiens AAA1 protein (AAA1), transcript variant VI, mRNA
 NM_207289 Homo sapiens AAA1 protein (AAA1), transcript variant VII, mRNA
 NM_207290 Homo sapiens AAA1 protein (AAA1), transcript variant VIII, mRNA
 NM_207291 Homo sapiens upstream transcription factor 2, c-fos interacting (USF2), tran
 NM_207292 Homo sapiens muscleblind-like (Drosophila) (MBNL1), transcript variant 2, m
 NM_207293 Homo sapiens muscleblind-like (Drosophila) (MBNL1), transcript variant 3, m
 NM_207294 Homo sapiens muscleblind-like (Drosophila) (MBNL1), transcript variant 4, m
 NM_207295 Homo sapiens muscleblind-like (Drosophila) (MBNL1), transcript variant 5, m
 NM_207296 Homo sapiens muscleblind-like (Drosophila) (MBNL1), transcript variant 6, m
 NM_207297 Homo sapiens muscleblind-like (Drosophila) (MBNL1), transcript variant 7, m
 NM_207299 Homo sapiens plasticity related gene 3 (PRG-3), transcript variant 1, mRNA
 NM_207303 Homo sapiens attractin-like 1 (ATRNL1), mRNA
 NM_207304 Homo sapiens muscleblind-like 2 (Drosophila) (MBNL2), transcript variant 3,
 NM_207305 Homo sapiens forkhead box D4 (FOXO4), mRNA
 NM_207306 Homo sapiens KIAA0495 (KIAA0495), mRNA
 NM_207307 Homo sapiens hypothetical protein LOC90288 (LOC90288), mRNA
 NM_207308 Homo sapiens nuclear pore membrane glycoprotein 210-like (LOC91181), m
 NM_207309 Homo sapiens UDP-N-acetylglucosamine pyrophosphorylase 1-like 1 (UAP1
 NM_207310 Homo sapiens hypothetical protein DKFZp434E2321 (DKFZp434E2321), mF
 NM_207311 Homo sapiens hypothetical protein LOC92558 (LOC92558), mRNA
 NM_207312 Homo sapiens similar to alpha tubulin (LOC112714), mRNA
 NM_207313 Homo sapiens hypothetical protein LOC124842 (LOC124842), mRNA
 NM_207314 Homo sapiens VNFT9373 (UNQ9373), mRNA
 NM_207315 Homo sapiens hypothetical protein LOC129607 (LOC129607), mRNA

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NM_207316 Homo sapiens SRSR846 (UNQ846), mRNA
 NM_207317 Homo sapiens FLJ32921 protein (FLJ32921), mRNA
 NM_207319 Homo sapiens FLJ32867 protein (FLJ32867), mRNA
 NM_207320 Homo sapiens FLJ25831 protein (FLJ25831), mRNA
 NM_207321 Homo sapiens chromosome 10 open reading frame 129 (C10orf129), mRNA
 NM_207322 Homo sapiens hypothetical LOC145741 (LOC145741), mRNA
 NM_207323 Homo sapiens hypothetical protein DKFZp667M2411 (DKFZp667M2411), mRNA
 NM_207324 Homo sapiens hypothetical protein LOC147650 (LOC147650), mRNA
 NM_207325 Homo sapiens hypothetical protein LOC147991 (LOC147991), mRNA
 NM_207326 Homo sapiens hypothetical protein LOC149134 (LOC149134), mRNA
 NM_207327 Homo sapiens similar to RIKEN cDNA 2210021J22 (LOC150383), mRNA
 NM_207328 Homo sapiens hypothetical protein LOC150763 (LOC150763), mRNA
 NM_207329 Homo sapiens myeloid-associated differentiation marker-like (MYADML), mRNA
 NM_207330 Homo sapiens hypothetical protein LOC152519 (LOC152519), mRNA
 NM_207331 Homo sapiens hypothetical protein LOC153561 (LOC153561), mRNA
 NM_207332 Homo sapiens hypothetical protein LOC157697 (LOC157697), mRNA
 NM_207333 Homo sapiens hypothetical protein LOC162967 (LOC162967), mRNA
 NM_207334 Homo sapiens family with sequence similarity 43, member B (FAM43B), mRNA
 NM_207335 Homo sapiens FLJ46299 protein (FLJ46299), mRNA
 NM_207336 Homo sapiens likely ortholog of mouse zinc finger protein EZI (EZI), mRNA
 NM_207337 Homo sapiens hypothetical protein LOC196394 (LOC196394), mRNA
 NM_207338 Homo sapiens likely ortholog of mouse klotho lactase-phlorizin hydrolase rek
 NM_207339 Homo sapiens similar to PAGE-5 protein (MGC62094), mRNA
 NM_207340 Homo sapiens hypothetical protein LOC254359 (LOC254359), mRNA
 NM_207341 Homo sapiens similar to ZP1 precursor (MGC87693), mRNA
 NM_207343 Homo sapiens hypothetical protein DKFZp547C195 (DKFZp547C195), mRNA
 NM_207344 Homo sapiens hypothetical protein LOC283377 (LOC283377), mRNA
 NM_207345 Homo sapiens HEEB9341 (UNQ9341), mRNA
 NM_207346 Homo sapiens likely homolog of yeast SEN54 (SEN54L), mRNA
 NM_207347 Homo sapiens chromosome 18 open reading frame 30 (C18orf30), mRNA
 NM_207348 Homo sapiens hypothetical protein LOC284723 (LOC284723), mRNA
 NM_207349 Homo sapiens hypothetical protein LOC284739 (LOC284739), mRNA
 NM_207350 Homo sapiens similar to FRG1 protein (FSHD region gene 1 protein) (MGC7.
 NM_207351 Homo sapiens hypothetical protein FLJ33674 (FLJ33674), mRNA
 NM_207352 Homo sapiens cytochrome P450, family 4, subfamily V, polypeptide 2 (CYP4
 NM_207353 Homo sapiens ubiquitin-conjugating enzyme UbcM2 (LOC286480), mRNA
 NM_207354 Homo sapiens hypothetical protein LOC338692 (LOC338692), mRNA
 NM_207355 Homo sapiens protein expressed in prostate, ovary, testis, and placenta (PO'
 NM_207356 Homo sapiens hypothetical protein LOC339448 (LOC339448), mRNA
 NM_207357 Homo sapiens hypothetical protein LOC339524 (LOC339524), mRNA
 NM_207358 Homo sapiens hypothetical protein LOC339789 (LOC339789), mRNA
 NM_207359 Homo sapiens glutamate decarboxylase-like 1 (GADL1), mRNA
 NM_207362 Homo sapiens similar to 2010300C02Rik protein (MGC42367), mRNA
 NM_207363 Homo sapiens Nck-associated protein 5 (NAP5), mRNA
 NM_207364 Homo sapiens G protein-coupled receptor 148 (GPR148), mRNA
 NM_207365 Homo sapiens similar to Arylacetamide deacetylase (AADAC) (MGC72001),
 NM_207366 Homo sapiens FLJ44060 protein (FLJ44060), mRNA
 NM_207367 Homo sapiens FLJ42291 protein (FLJ42291), mRNA
 NM_207368 Homo sapiens hypothetical protein LOC348262 (LOC348262), mRNA
 NM_207370 Homo sapiens G protein-coupled receptor 153 (GPR153), mRNA
 NM_207371 Homo sapiens FLJ45187 protein (FLJ45187), mRNA
 NM_207372 Homo sapiens SH2 domain containing 4B (SH2D4B), mRNA
 NM_207373 Homo sapiens chromosome 10 open reading frame 99 (C10orf99), mRNA
 NM_207374 Homo sapiens olfactory receptor (UNQ6466), mRNA
 NM_207375 Homo sapiens INPE5792 (UNQ5792), mRNA
 NM_207376 Homo sapiens hypothetical protein (LOC387882), mRNA
 NM_207377 Homo sapiens TIMM9 (UNQ9438), mRNA

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NM_207378 Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1 anti
 NM_207379 Homo sapiens FLJ42486 protein (FLJ42486), mRNA
 NM_207380 Homo sapiens FLJ43339 protein (FLJ43339), mRNA
 NM_207381 Homo sapiens FLJ41287 protein (FLJ41287), mRNA
 NM_207382 Homo sapiens FLJ43276 protein (FLJ43276), mRNA
 NM_207383 Homo sapiens FLJ42289 protein (FLJ42289), mRNA
 NM_207384 Homo sapiens QRWT5810 (UNQ5810), mRNA
 NM_207385 Homo sapiens FLJ26184 protein (FLJ26184), mRNA
 NM_207386 Homo sapiens FLJ45455 protein (FLJ45455), mRNA
 NM_207387 Homo sapiens FLJ35696 protein (FLJ35696), mRNA
 NM_207388 Homo sapiens FLJ31222 protein (FLJ31222), mRNA
 NM_207389 Homo sapiens FLJ44861 protein (FLJ44861), mRNA
 NM_207390 Homo sapiens FLJ45910 protein (FLJ45910), mRNA
 NM_207391 Homo sapiens FLJ45744 protein (FLJ45744), mRNA
 NM_207392 Homo sapiens KIPV467 (UNQ467), mRNA
 NM_207393 Homo sapiens insulin growth factor-like family member 3 (IGFL3), mRNA
 NM_207394 Homo sapiens FLJ45949 protein (FLJ45949), mRNA
 NM_207395 Homo sapiens FLJ45850 protein (FLJ45850), mRNA
 NM_207396 Homo sapiens FLJ46380 protein (FLJ46380), mRNA
 NM_207397 Homo sapiens EAPG6122 (UNQ6122), mRNA
 NM_207398 Homo sapiens FLJ38822 protein (FLJ38822), mRNA
 NM_207399 Homo sapiens FLJ36116 protein (FLJ36116), mRNA
 NM_207400 Homo sapiens FLJ39739 protein (FLJ39739), mRNA
 NM_207401 Homo sapiens FLJ45717 protein (FLJ45717), mRNA
 NM_207402 Homo sapiens FLJ42986 protein (FLJ42986), mRNA
 NM_207403 Homo sapiens FLJ45880 protein (FLJ45880), mRNA
 NM_207404 Homo sapiens FLJ46481 protein (FLJ46481), mRNA
 NM_207405 Homo sapiens FLJ43965 protein (FLJ43965), mRNA
 NM_207406 Homo sapiens FLJ16046 protein (FLJ16046), mRNA
 NM_207407 Homo sapiens FLJ27505 protein (FLJ27505), mRNA
 NM_207408 Homo sapiens AAAL3045 (UNQ3045), mRNA
 NM_207409 Homo sapiens IVF19356 (UNQ9356), mRNA
 NM_207410 Homo sapiens HARL2754 (UNQ2754), mRNA
 NM_207411 Homo sapiens FLJ43582 protein (FLJ43582), mRNA
 NM_207412 Homo sapiens RPLK9433 (UNQ9433), mRNA
 NM_207413 Homo sapiens FLJ43860 protein (FLJ43860), mRNA
 NM_207414 Homo sapiens FLJ44082 protein (FLJ44082), mRNA
 NM_207415 Homo sapiens FLJ46082 protein (FLJ46082), mRNA
 NM_207416 Homo sapiens Similar to RIKEN cDNA 2700049P18 gene (MGC57827), mR
 NM_207417 Homo sapiens C1q and tumor necrosis factor related protein 8 (C1QTNF8), r
 NM_207418 Homo sapiens Intestinal facilitative glucose transporter 7 (SLC2A7), mRNA
 NM_207419 Homo sapiens peptidylarginine deiminase type 6 (PADI6), mRNA
 NM_207420 Homo sapiens FLJ44635 protein (FLJ44635), mRNA
 NM_207421 Homo sapiens FLJ45983 protein (FLJ45983), mRNA
 NM_207422 Homo sapiens FLJ40536 protein (FLJ40536), mRNA
 NM_207423 Homo sapiens FLJ46831 protein (FLJ46831), mRNA
 NM_207424 Homo sapiens hypothetical gene supported by AY129010 (LOC399851), mR
 NM_207425 Homo sapiens FLJ45212 protein (FLJ45212), mRNA
 NM_207426 Homo sapiens FLJ45803 protein (FLJ45803), mRNA
 NM_207427 Homo sapiens FLJ46266 protein (FLJ46266), mRNA
 NM_207428 Homo sapiens FLJ45436 protein (FLJ45436), mRNA
 NM_207429 Homo sapiens FLJ44874 protein (FLJ44874), mRNA
 NM_207430 Homo sapiens FLJ46363 protein (FLJ46363), mRNA
 NM_207431 Homo sapiens FLJ40142 protein (FLJ40142), mRNA
 NM_207432 Homo sapiens FLJ42957 protein (FLJ42957), mRNA
 NM_207433 Homo sapiens FLJ43486 protein (FLJ43486), mRNA
 NM_207434 Homo sapiens FLJ43808 protein (FLJ43808), mRNA

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NM_207439 Homo sapiens FLJ46358 protein (FLJ46358), mRNA
NM_207440 Homo sapiens FLJ26443 protein (FLJ26443), mRNA
NM_207441 Homo sapiens FLJ42220 protein (FLJ42220), mRNA
NM_207442 Homo sapiens FLJ39779 protein (FLJ39779), mRNA
NM_207443 Homo sapiens FLJ45244 protein (FLJ45244), mRNA
NM_207444 Homo sapiens FLJ35895 protein (FLJ35895), mRNA
NM_207445 Homo sapiens FLJ39531 protein (FLJ39531), mRNA
NM_207446 Homo sapiens hypothetical gene supported by AK075564; BC060873 (LOC4
NM_207447 Homo sapiens IFM09370 (UNQ9370), mRNA
NM_207448 Homo sapiens FLJ45256 protein (FLJ45256), mRNA
NM_207449 Homo sapiens FLJ44674 protein (FLJ44674), mRNA
NM_207450 Homo sapiens FLJ27243 protein (FLJ27243), mRNA
NM_207451 Homo sapiens FLJ45121 protein (FLJ45121), mRNA
NM_207452 Homo sapiens FLJ45200 protein (FLJ45200), mRNA
NM_207453 Homo sapiens FLJ35934 protein (FLJ35934), mRNA
NM_207454 Homo sapiens FLJ44815 protein (FLJ44815), mRNA
NM_207458 Homo sapiens FLJ46026 protein (FLJ46026), mRNA
NM_207459 Homo sapiens FLJ35767 protein (FLJ35767), mRNA
NM_207460 Homo sapiens FLJ44313 protein (FLJ44313), mRNA
NM_207461 Homo sapiens FLJ44881 protein (FLJ44881), mRNA
NM_207462 Homo sapiens FLJ45684 protein (FLJ45684), mRNA
NM_207463 Homo sapiens FLJ46230 protein (FLJ46230), mRNA
NM_207464 Homo sapiens FLJ40008 protein (FLJ40008), mRNA
NM_207465 Homo sapiens FLJ45337 protein (FLJ45337), mRNA
NM_207466 Homo sapiens FLJ46489 protein (FLJ46489), mRNA
NM_207467 Homo sapiens FLJ35530 protein (FLJ35530), mRNA
NM_207468 Homo sapiens FLJ43505 protein (FLJ43505), mRNA
NM_207469 Homo sapiens KFL1827 (UNQ827), mRNA
NM_207470 Homo sapiens FLJ45832 protein (FLJ45832), mRNA
NM_207471 Homo sapiens FLJ42200 protein (FLJ42200), mRNA
NM_207472 Homo sapiens FLJ46020 protein (FLJ46020), mRNA
NM_207473 Homo sapiens FLJ41733 protein (FLJ41733), mRNA
NM_207474 Homo sapiens FLJ42953 protein (FLJ42953), mRNA
NM_207475 Homo sapiens FLJ90680 protein (FLJ90680), mRNA
NM_207477 Homo sapiens FLJ27365 protein (FLJ27365), mRNA
NM_207478 Homo sapiens FLJ44385 protein (FLJ44385), mRNA
NM_207479 Homo sapiens FLJ41046 protein (FLJ41046), mRNA
NM_207480 Homo sapiens AILT5830 (UNQ5830), mRNA
NM_207481 Homo sapiens FLJ34870 protein (FLJ34870), mRNA
NM_207482 Homo sapiens FLJ44048 protein (FLJ44048), mRNA
NM_207483 Homo sapiens FLJ45964 protein (FLJ45964), mRNA
NM_207484 Homo sapiens FLJ40712 protein (FLJ40712), mRNA
NM_207485 Homo sapiens FLJ41327 protein (FLJ41327), mRNA
NM_207486 Homo sapiens FLJ44076 protein (FLJ44076), mRNA
NM_207487 Homo sapiens FLJ46211 protein (FLJ46211), mRNA
NM_207488 Homo sapiens FLJ42393 protein (FLJ42393), mRNA
NM_207489 Homo sapiens FLJ35816 protein (FLJ35816), mRNA
NM_207490 Homo sapiens FLJ45721 protein (FLJ45721), mRNA
NM_207491 Homo sapiens similar to KIAA1680 protein (MGC48628), mRNA
NM_207492 Homo sapiens FLJ44477 protein (FLJ44477), mRNA
NM_207493 Homo sapiens FLJ44896 protein (FLJ44896), mRNA
NM_207494 Homo sapiens FLJ40092 protein (FLJ40092), mRNA
NM_207495 Homo sapiens hypothetical protein DKFzp686i15217 (DKFzp686i15217), ml
NM_207496 Homo sapiens chromosome 6 open reading frame 214 (C6orf214), mRNA
NM_207497 Homo sapiens FLJ43752 protein (FLJ43752), mRNA
NM_207498 Homo sapiens FLJ43093 protein (FLJ43093), mRNA
NM_207499 Homo sapiens FLJ41841 protein (FLJ41841), mRNA

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NM_207500 Homo sapiens FLJ44955 protein (FLJ44955), mRNA
 NM_207501 Homo sapiens FLJ27255 protein (FLJ27255), mRNA
 NM_207502 Homo sapiens chromosome 6 open reading frame 122 (C6orf122), mRNA
 NM_207503 Homo sapiens FLJ42280 protein (FLJ42280), mRNA
 NM_207504 Homo sapiens FLJ46365 protein (FLJ46365), mRNA
 NM_207505 Homo sapiens FLJ45248 protein (FLJ45248), mRNA
 NM_207506 Homo sapiens sterile alpha motif domain containing 12 (SAMD12), mRNA
 NM_207507 Homo sapiens FLJ45202 protein (FLJ45202), mRNA
 NM_207508 Homo sapiens FLJ45478 protein (FLJ45478), mRNA
 NM_207509 Homo sapiens FLJ46836 protein (FLJ46836), mRNA
 NM_207510 Homo sapiens FLJ45224 protein (FLJ45224), mRNA
 NM_207511 Homo sapiens FLJ36268 protein (FLJ36268), mRNA
 NM_207512 Homo sapiens nuclear RNA export factor-like (LOC401610), mRNA
 NM_207513 Homo sapiens POTE14 (LOC404785), mRNA
 NM_207514 Homo sapiens hypothetical protein FLJ20186 (FLJ20186), transcript variant 1
 NM_207517 Homo sapiens ADAMTS-like 3 (ADAMTSL3), mRNA
 NM_207518 Homo sapiens protein kinase, cAMP-dependent, catalytic, alpha (PRKACA), mRNA
 NM_207519 Homo sapiens zeta-chain (TCR) associated protein kinase 70kDa (ZAP70), mRNA
 NM_207520 Homo sapiens reticulin 4 (RTN4), transcript variant 4, mRNA
 NM_207521 Homo sapiens reticulin 4 (RTN4), transcript variant 5, mRNA
 NM_207577 Homo sapiens microtubule-associated protein 6 (MAP6), transcript variant 2, mRNA
 NM_207578 Homo sapiens protein kinase, cAMP-dependent, catalytic, beta (PRKACB), mRNA
 NM_207581 Homo sapiens similar to Numb-interacting homolog gene (LOC405753), mRNA
 NM_207582 Homo sapiens HERV-FRD provirus ancestral Env polyprotein (HERV-FRD), mRNA
 NM_207584 Homo sapiens interferon (alpha, beta and omega) receptor 2 (IFNAR2), mRNA
 NM_207585 Homo sapiens Interferon (alpha, beta and omega) receptor 2 (IFNAR2), transcript variant 1
 NM_207627 Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 1 (ABCG1), mRNA
 NM_207628 Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 1 (ABCG1), mRNA
 NM_207629 Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 1 (ABCG1), mRNA
 NM_207630 Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 1 (ABCG1), mRNA
 NM_207644 Homo sapiens similar to hypothetical protein LOC192734 (LOC388886), mRNA
 NM_207645 Homo sapiens similar to expressed sequence A1593442 (LOC399947), mRNA
 NM_207646 Homo sapiens eosinophil lysophospholipase-like (LOC400696), mRNA
 NM_207647 Homo sapiens similar to fibronectin type 3 and SPRY domain-containing protein 1
 NM_207660 Homo sapiens nuclear protein UKp68 (FLJ11806), transcript variant 2, mRNA
 NM_207661 Homo sapiens nuclear protein UKp68 (FLJ11806), transcript variant 3, mRNA
 NM_207662 Homo sapiens nuclear protein UKp68 (FLJ11806), transcript variant 4, mRNA
 NM_207672 Homo sapiens GRIP1 associated protein 1 (GRIPAP1), transcript variant 2, mRNA
 NM_212460 Homo sapiens ADP-ribosylation factor-like 4A (ARL4A), transcript variant 2, mRNA
 NM_212461 Homo sapiens protein kinase, AMP-activated, gamma 1 non-catalytic subunit 1
 NM_212464 Homo sapiens calpain 3, (p94) (CAPN3), transcript variant 8, mRNA
 NM_212465 Homo sapiens calpain 3, (p94) (CAPN3), transcript variant 7, mRNA
 NM_212467 Homo sapiens calpain 3, (p94) (CAPN3), transcript variant 9, mRNA
 NM_212469 Homo sapiens choline kinase alpha (CHKa), transcript variant 2, mRNA
 NM_212471 Homo sapiens protein kinase, cAMP-dependent, regulatory, type I, alpha (ti) (PRKAR1A), mRNA
 NM_212472 Homo sapiens protein kinase, cAMP-dependent, regulatory, type I, alpha (ti) (PRKAR1A), mRNA
 NM_212474 Homo sapiens fibronectin 1 (FN1), transcript variant 6, mRNA
 NM_212475 Homo sapiens fibronectin 1 (FN1), transcript variant 2, mRNA
 NM_212476 Homo sapiens fibronectin 1 (FN1), transcript variant 5, mRNA
 NM_212478 Homo sapiens fibronectin 1 (FN1), transcript variant 4, mRNA
 NM_212479 Homo sapiens zinc finger, MYND domain containing 11 (ZMYND11), transcript variant 1
 NM_212481 Homo sapiens AT rich interactive domain 5A (MRP1-like) (ARID5A), transcript variant 1
 NM_212482 Homo sapiens fibronectin 1 (FN1), transcript variant 1, mRNA
 NM_212492 Homo sapiens G protein pathway suppressor 1 (GPS1), transcript variant 1, mRNA
 NM_212502 Homo sapiens PCTAIRE protein kinase 3 (PCTK3), transcript variant 2, mRNA
 NM_212503 Homo sapiens PCTAIRE protein kinase 3 (PCTK3), transcript variant 1, mRNA
 NM_212530 Homo sapiens cell division cycle 25B (CDC25B), transcript variant 5, mRNA

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NM_212533 Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 2 (ABC2)
 NM_212535 Homo sapiens protein kinase C, beta 1 (PRKCB1), transcript variant 1, mRNA
 NM_212539 Homo sapiens protein kinase C, delta (PRKCD), transcript variant 2, mRNA
 NM_212540 Homo sapiens E2F transcription factor 6 (E2F6), transcript variant f, mRNA
 NM_212543 Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide
 NM_212550 Homo sapiens biogenesis of lysosome-related organelles complex-1, subunit
 NM_212551 Homo sapiens hypothetical protein SB145 (SB145), mRNA
 NM_212552 Homo sapiens similar to RIKEN 1810056O20 (LOC388962), mRNA
 NM_212553 Homo sapiens deubiquitinating enzyme DUB4 (DUB4), mRNA
 NM_212554 Homo sapiens similar to CG9643-PA (LOC399818), mRNA
 NM_212555 Homo sapiens LVL3112 (UNQ3112), mRNA
 NM_212556 Homo sapiens ankyrin repeat and SOCS box-containing 18 (ASB18), mRNA
 NM_212557 Homo sapiens RST1889 (UNQ689), mRNA
 NM_212558 Homo sapiens similar to RIKEN A930001M12 (LOC401498), mRNA
 NM_212559 Homo sapiens X Kell blood group precursor-related, X-linked (XKRX), mRNA
 NM_213590 Homo sapiens protein kinase N1 (PKN1), transcript variant 1, mRNA
 NM_213596 Homo sapiens DNA fragmentation factor, 45kDa, alpha polypeptide (DFFA),
 NM_213598 Homo sapiens solute carrier family 39 (zinc transporter), member 3 (SLC39A3)
 NM_213599 Homo sapiens nebulin (NEBL), transcript variant 2, mRNA
 NM_213599 Homo sapiens Ras association (RalGDS/AF-6) and pleckstrin homology domain
 NM_213590 Homo sapiens ret finger protein 2 (RFP2), transcript variant 3, mRNA
 NM_213593 Homo sapiens deiodinase, iodothyronine, type I (DIO1), transcript variant 2, mRNA
 NM_213594 Homo sapiens regulatory factor X, 4 (influences HLA class II expression) (RF)
 NM_213596 Homo sapiens forkhead box N4 (FOXN4), mRNA
 NM_213597 Homo sapiens hypothetical protein LOC124751 (LOC124751), mRNA
 NM_213598 Homo sapiens zinc finger protein 543 (ZNF543), mRNA
 NM_213599 Homo sapiens transmembrane protein 16E (TMEM16E), mRNA
 NM_213600 Homo sapiens hypothetical protein LOC255189 (LOC255189), mRNA
 NM_213601 Homo sapiens hypothetical protein LOC283578 (LOC283578), mRNA
 NM_213602 Homo sapiens CD33 antigen-like 3 (CD33L3), mRNA
 NM_213603 Homo sapiens hypothetical protein LOC285989 (LOC285989), mRNA
 NM_213604 Homo sapiens thrombospondin, type I, domain containing 6 (THSD6), mRNA
 NM_213605 Homo sapiens zinc finger protein 517 (ZNF517), mRNA
 NM_213606 Homo sapiens similar monocarboxylate transporter (LOC387700), mRNA
 NM_213607 Homo sapiens similar to RIKEN 4933439F11 (LOC388389), mRNA
 NM_213608 Homo sapiens ILDS411 (UNQ6411), mRNA
 NM_213609 Homo sapiens TAF11 protein (TAF11), mRNA
 NM_213611 Homo sapiens solute carrier family 25 (mitochondrial carrier; phosphate carrier)
 NM_213612 Homo sapiens solute carrier family 25 (mitochondrial carrier; phosphate carrier)
 NM_213613 Homo sapiens solute carrier family 26 (sulfate transporter), member 1 (SLC26)
 NM_213618 Homo sapiens suppression of tumorigenicity 5 (ST5), transcript variant 3, mRNA
 NM_213619 Homo sapiens ATPase, H⁺ transporting, lysosomal 50/57kDa, V1 subunit H
 NM_213620 Homo sapiens ATPase, H⁺ transporting, lysosomal 50/57kDa, V1 subunit H
 NM_213621 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 3A (HTR3A), transcript
 NM_213622 Homo sapiens STAM binding protein (STAMPB), transcript variant 3, mRNA
 NM_213631 Homo sapiens chromosome 20 open reading frame 132 (C20orf132), transcript
 NM_213632 Homo sapiens chromosome 20 open reading frame 132 (C20orf132), transcript
 NM_213633 Homo sapiens pregnancy specific beta-1-glycoprotein 4 (PSG4), transcript variant
 NM_213636 Homo sapiens PDZ and LIM domain 7 (enigma) (PDLIM7), transcript variant
 NM_213645 Homo sapiens tryptophanyl-tRNA synthetase (WARS), transcript variant 3, mRNA
 NM_213646 Homo sapiens tryptophanyl-tRNA synthetase (WARS), transcript variant 4, mRNA
 NM_213647 Homo sapiens fibroblast growth factor receptor 4 (FGFR4), transcript variant
 NM_213648 Homo sapiens transcription factor 7 (T-cell specific, HMG-box) (TCF7), transcript
 NM_213649 Homo sapiens sideroflexin 4 (SFXN4), transcript variant 1, mRNA
 NM_213650 Homo sapiens sideroflexin 4 (SFXN4), transcript variant 3, mRNA
 NM_213651 Homo sapiens solute carrier family 25 (mitochondrial carrier; phosphate carrier)
 NM_213652 Homo sapiens hemochromatosis type 2 (juvenile) (HFE2), transcript variant c

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NM_213653 Homo sapiens hemochromatosis type 2 (juvenile) (HFE2), transcript variant 1
 NM_213654 Homo sapiens armadillo repeat containing 8 (ARMC8), mRNA
 NM_213655 Homo sapiens hereditary sensory neuropathy, type II (HSN2), mRNA
 NM_213656 Homo sapiens type I hair keratin KA35 (KA35), mRNA
 NM_213657 Homo sapiens killer cell lectin-like receptor subfamily C, member 1 (KLCRC1),
 NM_213658 Homo sapiens killer cell lectin-like receptor subfamily C, member 1 (KLCRC1),
 NM_213662 Homo sapiens signal transducer and activator of transcription 3 (acute-phase
 NM_213674 Homo sapiens tropomyosin 2 (beta) (TPM2), transcript variant 2, mRNA
 NM_213720 Homo sapiens chromosome 22 open reading frame 16 (C22orf16), mRNA
 NM_213723 Homo sapiens chromosome 13 open reading frame 25 (C13orf25), transcript
 NM_213724 Homo sapiens chromosome 13 open reading frame 25 (C13orf25), transcript
 NM_213725 Homo sapiens ribosomal protein, large, P1 (RPLP1), transcript variant 2, mR
 NM_213726 Homo sapiens inhibitor of CDK interacting with cyclin A1 (INCA1), mRNA
 NM_214461 Homo sapiens MGC50273 protein (MGC50273), mRNA
 NM_214462 Homo sapiens dapper homolog 2, antagonist of beta-catenin (xenopus) (DAP
 NM_214675 Homo sapiens CD209 antigen-like (CD209L), transcript variant 2, mRNA
 NM_214676 Homo sapiens CD209 antigen-like (CD209L), transcript variant 3, mRNA
 NM_214677 Homo sapiens CD209 antigen-like (CD209L), transcript variant 4, mRNA
 NM_214678 Homo sapiens CD209 antigen-like (CD209L), transcript variant 5, mRNA
 NM_214679 Homo sapiens CD209 antigen-like (CD209L), transcript variant 6, mRNA
 NM_214710 Homo sapiens protease, serine-like 1 (PRSSL1), mRNA
 NM_214711 Homo sapiens hypothetical LOC401137 (LOC401137), mRNA
 XM_001279 Homo sapiens phosphoprotein enriched in astrocytes 15 (PEA15), mRNA
 XM_001290 Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 4 (ABC4
 XM_001296 Homo sapiens cytosolic acyl coenzyme A thioester hydrolase (hBACH), mR
 XM_001322 Homo sapiens coagulation factor III (thromboplastin, tissue factor) (F3), mR
 XM_001393 Homo sapiens peroxiredoxin 1 (PRDX1), mRNA
 XM_001442 Homo sapiens phosphoglucomutase 1 (PGM1), mRNA
 XM_001463 Homo sapiens ATPase, H⁺ transporting, lysosomal (vacuolar proton pump) 2
 XM_001527 Homo sapiens polymyositis/scleroderma autoantigen 2 (100kD) (PMSCL2), r
 XM_001541 Homo sapiens heterogeneous nuclear ribonucleoprotein R (HNRPR), mRNA
 XM_001607 Homo sapiens growth arrest and DNA-damage-inducible, alpha (GADD45A),
 XM_001644 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 8 (PPP1f
 XM_001654 Homo sapiens chromosome 1 open reading frame 8 (C1orf8), mRNA
 XM_001655 Homo sapiens HSPCO34 protein (LOC51668), mRNA
 XM_001677 Homo sapiens involucrin (IVL), mRNA
 XM_001690 Homo sapiens cytochrome b5 reductase 1 (B5R.1) (LOC51706), mRNA
 XM_007651 Homo sapiens similar to Sorbitol dehydrogenase (L-iditol 2-dehydrogenase) 1
 XM_010658 Homo sapiens similar to protein phosphatase 1, regulatory (inhibitor) subunit
 XM_012219 Homo sapiens similar to Phosphoglycerate mutase 1 (Phosphoglycerate mu
 XM_015334 Homo sapiens family with sequence similarity 10, member A3 (FAM10A3), m
 XM_015717 Homo sapiens similar to 40S ribosomal protein S7 (S8) (LOC149224), mRN
 XM_016093 Homo sapiens similar to eukaryotic initiation factor 5A isoform 1 variant A (LC
 XM_016113 Homo sapiens similar to 40S ribosomal protein S10 (LOC158104), mRNA
 XM_016532 Homo sapiens similar to hepatitis C virus core-binding protein 6; cervical can
 XM_016548 Homo sapiens chromodomain protein, Y chromosome, 2 related (LOC20361
 XM_016713 Homo sapiens similar to ribosomal protein S3a; 40S ribosomal protein S3a; 1
 XM_017374 Homo sapiens similar to Nonhistone chromosomal protein HMG-17 (H-high-mc
 XM_017681 Homo sapiens similar to 40S ribosomal protein S26 (LOC158200), mRNA
 XM_017966 Homo sapiens similar to Reticulon protein 3 (Neuroendocrine-specific prot
 XM_018399 Homo sapiens hypothetical protein LOC144983 (LOC144983), mRNA
 XM_018432 Homo sapiens similar to 60S ribosomal protein L7 (LOC146110), mRNA
 XM_018487 Homo sapiens similar to omega protein (LOC91353), mRNA
 XM_027045 Homo sapiens cut-like 2 (Drosophila) (CUTL2), mRNA
 XM_027074 Homo sapiens (3)mbt-like 3 (Drosophila) (L3MBTL3), mRNA
 XM_027105 Homo sapiens KIAA0767 protein (KIAA0767), mRNA
 XM_027162 Homo sapiens DMRT-like family A2 (DMRTA2), mRNA

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XM_027236 Homo sapiens tetra-tryptophan repeat domain 9 (TTC9), mRNA
 XM_027237 Homo sapiens mitogen-activated protein kinase kinase 9 (MAP3K9), mRNA
 XM_027307 Homo sapiens pleckstrin homology domain containing, family G (with RhoGEF), mRNA
 XM_027330 Homo sapiens RNA-binding region (RNP1, RRM) containing 7 (RNPC7), mRNA
 XM_027658 Homo sapiens fibronectin type III domain containing 1 (FNDC1), mRNA
 XM_028067 Homo sapiens midline (MIDN), mRNA
 XM_028217 Homo sapiens hypothetical LOC90024 (LOC90024), mRNA
 XM_028253 Homo sapiens chromosome 19 open reading frame 7 (C19orf7), mRNA
 XM_028413 Homo sapiens KIAA1374 protein (KIAA1374), mRNA
 XM_028522 Homo sapiens myosin heavy chain Myr 8 (MYR8), mRNA
 XM_028810 Homo sapiens KIAA1755 protein (KIAA1755), mRNA
 XM_029084 Homo sapiens hypothetical protein FLJ21438 (FLJ21438), mRNA
 XM_029101 Homo sapiens KIAA0947 protein (KIAA0947), mRNA
 XM_029323 Homo sapiens hypothetical protein LOC90133 (LOC90133), mRNA
 XM_029353 Homo sapiens KIAA1509 (KIAA1509), mRNA
 XM_029429 Homo sapiens KIAA1328 protein (KIAA1328), mRNA
 XM_029438 Homo sapiens KIAA0397 gene product (KIAA0397), mRNA
 XM_029805 Homo sapiens similar to ribosomal protein L7 (LOC90193), mRNA
 XM_029992 Homo sapiens potassium channel, subfamily T, member 1 (KCN1T), mRNA
 XM_030300 Homo sapiens netrin receptor Unc5h1 (KIAA1978), mRNA
 XM_030378 Homo sapiens zinc finger protein 527 (ZNF527), mRNA
 XM_030445 Homo sapiens chromosome 10 open reading frame 75 (C10orf75), mRNA
 XM_030559 Homo sapiens par-6 partitioning defective 6 homolog beta (C. elegans) (PAF6), mRNA
 XM_030577 Homo sapiens ATPase, Class II, type 9A (ATP9A), mRNA
 XM_030665 Homo sapiens KIAA1229 protein (KIAA1229), mRNA
 XM_030689 Homo sapiens hypothetical protein LOC90288 (LOC90288), mRNA
 XM_030729 Homo sapiens hypothetical protein DKFZp434I1117 (DKFZp434I1117), mRNA
 XM_030892 Homo sapiens hypothetical protein LOC90317 (LOC90317), mRNA
 XM_030893 Homo sapiens similar to ribosomal protein L37 (LOC147655), mRNA
 XM_030896 Homo sapiens hypothetical protein LOC90321 (LOC90321), mRNA
 XM_030958 Homo sapiens hypothetical protein LOC90333 (LOC90333), mRNA
 XM_031009 Homo sapiens similar to fer-1 like protein 3 (LOC90342), mRNA
 XM_031102 Homo sapiens WD repeat domain 22 (WDR22), mRNA
 XM_031104 Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylglucosaminyltransferase 4 (ZSWIM4), mRNA
 XM_031246 Homo sapiens roundsabout, axon guidance receptor, homolog 2 (Drosophila) (ROUNDAABOUT), mRNA
 XM_031342 Homo sapiens zinc finger, SWIM domain containing 4 (ZSWIM4), mRNA
 XM_031357 Homo sapiens KIAA0802 protein (KIAA0802), mRNA
 XM_031401 Homo sapiens EGF-like-domain, multiple 3 (EGFL3), mRNA
 XM_031553 Homo sapiens U2-associated SR140 protein (SR140), mRNA
 XM_031561 Homo sapiens TRAF4 associated factor 1 (FLJ14502), mRNA
 XM_031689 Homo sapiens MAX gene associated (MGA), mRNA
 XM_031706 Homo sapiens likely ortholog of mouse mitogen activated protein kinase binc (BINC), mRNA
 XM_031744 Homo sapiens START domain containing 9 (STARD9), mRNA
 XM_031975 Homo sapiens similar to Ribulose-phosphate 3-epimerase (Ribulose-5-phosphate 3-epimerase), mRNA
 XM_032059 Homo sapiens similar to BC37295_3 (LOC90485), mRNA
 XM_032181 Homo sapiens KIAA1233 protein (KIAA1233), mRNA
 XM_032278 Homo sapiens signal-induced proliferation-associated 1 like 3 (SIPA1L3), mRNA
 XM_032397 Homo sapiens DKFZP564I122 protein (DKFZP564I122), mRNA
 XM_032542 Homo sapiens FLJ41352 protein (FLJ41352), mRNA
 XM_032571 Homo sapiens KIAA0888 protein (KIAA0888), mRNA
 XM_032678 Homo sapiens hypothetical protein LOC90576 (LOC90576), mRNA
 XM_032693 Homo sapiens KIAA0420 gene product (KIAA0420), mRNA
 XM_032812 Homo sapiens similar to hypothetical protein (LOC388506), mRNA
 XM_032901 Homo sapiens KIAA0226 gene product (KIAA0226), mRNA
 XM_032945 Homo sapiens chromosome 21 open reading frame 25 (C21orf25), mRNA
 XM_032996 Homo sapiens KIAA0819 protein (KIAA0819), mRNA
 XM_032997 Homo sapiens flavoprotein oxidoreductase MICAL3 (MICAL3), mRNA

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XM_039169 Homo sapiens KIAA1276 protein (KIAA1276), mRNA
 XM_039218 Homo sapiens similar to ribosomal protein S2; 40S ribosomal protein S2 (LO
 XM_039385 Homo sapiens KIAA1093 protein (KIAA1093), mRNA
 XM_039393 Homo sapiens plexin A4 (PLXNA4), mRNA
 XM_039495 Homo sapiens DNA segment, Chr 15, Wayne State University 75, expressed
 XM_039515 Homo sapiens G2 protein (G2), mRNA
 XM_039548 Homo sapiens SMYD family member 5 (SMYD5), mRNA
 XM_039570 Homo sapiens SEC15-like 2 (S. cerevisiae) (SEC15L2), mRNA
 XM_039627 Homo sapiens contactin 3 (plasmacytoma associated) (CNTN3), mRNA
 XM_039676 Homo sapiens KIAA1240 protein (KIAA1240), mRNA
 XM_039698 Homo sapiens KIAA1432 (KIAA1432), mRNA
 XM_039702 Homo sapiens similar to ribosomal protein S3a; 40S ribosomal protein S3a; 1
 XM_039721 Homo sapiens similar to MGC5244 protein (LOC91632), mRNA
 XM_039733 Homo sapiens KIAA0953 (KIAA0953), mRNA
 XM_039762 Homo sapiens myelin transcription factor 1-like (MYT1L), mRNA
 XM_039796 Homo sapiens Traf2 and NCK interacting kinase (KIAA0551), mRNA
 XM_039877 Homo sapiens mucin 5, subtype B, tracheobronchial (MUC5B), mRNA
 XM_039908 Homo sapiens hypothetical protein BC007307 (LOC91664), mRNA
 XM_039922 Homo sapiens similar to FLJ00050 protein (LOC401873), mRNA
 XM_040149 Homo sapiens similar to E74-like factor 2 (ets domain transcription factor); n
 XM_040265 Homo sapiens KIAA0217 protein (KIAA0217), mRNA
 XM_040383 Homo sapiens KIAA1677 (KIAA1677), mRNA
 XM_040486 Homo sapiens KIAA1789 protein (KIAA1789), mRNA
 XM_040527 Homo sapiens tenascin N (TNN), mRNA
 XM_040592 Homo sapiens zinc finger protein 469 (ZNF469), mRNA
 XM_040910 Homo sapiens chromosome 14 open reading frame 73 (C14orf73), mRNA
 XM_041018 Homo sapiens KIAA0367 (KIAA0367), mRNA
 XM_041020 Homo sapiens similar to protein 40kD (LOC158473), mRNA
 XM_041116 Homo sapiens chromosome 14 open reading frame 171 (C14orf171), mRNA
 XM_041126 Homo sapiens KIAA1486 protein (KIAA1486), mRNA
 XM_041162 Homo sapiens Nedd4 family interacting protein 2 (NDFIP2), mRNA
 XM_041191 Homo sapiens KIAA0931 protein (KIAA0931), mRNA
 XM_041221 Homo sapiens similar to RNA-binding protein S1, serine-rich domain; SR pro
 XM_041363 Homo sapiens likely ortholog of mouse semaF cytoplasmic domain associat
 XM_041964 Homo sapiens KIAA0523 protein (KIAA0523), mRNA
 XM_042068 Homo sapiens mitogen-activated protein kinase kinase 1 (MAP3K1), 1
 XM_042178 Homo sapiens similar to AKAP-binding sperm protein ropporin (LOC152015)
 XM_042234 Homo sapiens similar to RIKEN cDNA 4933437K13 (LOC92017), mRNA
 XM_042301 Homo sapiens KIAA1546 protein (KIAA1546), mRNA
 XM_042323 Homo sapiens calmodulin binding transcription activator 1 (CAMTA1), mRNA
 XM_042500 Homo sapiens similar to ribosomal protein S2; 40S ribosomal protein S2 (LO
 XM_042635 Homo sapiens KIAA1069 protein (KIAA1069), mRNA
 XM_042661 Homo sapiens KIAA1530 protein (KIAA1530), mRNA
 XM_042685 Homo sapiens KIAA1414 protein (KIAA1414), mRNA
 XM_042698 Homo sapiens ubiquitin specific protease 22 (USP22), mRNA
 XM_042833 Homo sapiens KIAA0295 protein (KIAA0295), mRNA
 XM_042936 Homo sapiens glutamate receptor interacting protein 2 (GRIP2), mRNA
 XM_042978 Homo sapiens KIAA1817 protein (KIAA1817), mRNA
 XM_043118 Homo sapiens KIAA0286 protein (KIAA0286), mRNA
 XM_043272 Homo sapiens KIAA0346 protein (KIAA0346), mRNA
 XM_043492 Homo sapiens KIAA1728 protein (KIAA1728), mRNA
 XM_043493 Homo sapiens synaptic vesicle protein 2C (SV2C), mRNA
 XM_043500 Homo sapiens similar to death-associated protein (LOC92196), mRNA
 XM_043613 Homo sapiens glutamate receptor, ionotropic, delta 1 (GRID1), mRNA
 XM_043624 Homo sapiens hypothetical protein DKFZp434E1822 (DKFZp434E1822), mF
 XM_043653 Homo sapiens hypothetical protein FLJ10097 (FLJ10097), mRNA
 XM_043739 Homo sapiens hypothetical cardiac/skeletal muscle-expressed ORF (LOC92

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XM_043863 Homo sapiens hypothetical protein DKFZp434H2226 (DKFZp434H2226), mf
 XM_043979 Homo sapiens similar to FLJ12363 protein (LOC92267), mRNA
 XM_043989 Homo sapiens hypothetical protein LOC92270 (LOC92270), mRNA
 XM_044062 Homo sapiens hypothetical protein DKFZp761O2018 (DKFZp761O2018), ml
 XM_044166 Homo sapiens hypothetical protein LOC92312 (LOC92312), mRNA
 XM_044178 Homo sapiens KIAA1211 protein (KIAA1211), mRNA
 XM_044196 Homo sapiens DKFZP434C212 protein (DKFZP434C212), mRNA
 XM_044212 Homo sapiens KIAA1862 protein (KIAA1862), mRNA
 XM_044334 Homo sapiens RIM binding protein 2 (KIAA0318), mRNA
 XM_044434 Homo sapiens KIAA1458 protein (KIAA1458), mRNA
 XM_044461 Homo sapiens KIAA1102 protein (KIAA1102), mRNA
 XM_044580 Homo sapiens KIAA1024 protein (KIAA1024), mRNA
 XM_044622 Homo sapiens collagen, type XIV, alpha 1 (undulin) (COL14A1), mRNA
 XM_044632 Homo sapiens KIAA0556 protein (KIAA0556), mRNA
 XM_044727 Homo sapiens myotubularin related protein 7 (MTMR7), mRNA
 XM_044836 Homo sapiens KIAA1340 protein (KIAA1340), mRNA
 XM_044921 Homo sapiens KIAA1442 protein (KIAA1442), mRNA
 XM_045086 Homo sapiens KIAA1764 protein (KIAA1764), mRNA
 XM_045113 Homo sapiens astrotactin (ASTN), mRNA
 XM_045271 Homo sapiens KIAA1580 protein (KIAA1580), mRNA
 XM_045283 Homo sapiens similar to IK cytokine; arginine/glutamic acid/aspartic acid rep
 XM_045290 Homo sapiens similar to basic leucine zipper and W2 domains 1 (LOC15157
 XM_045308 Homo sapiens PHD finger protein 19 (PHF19), mRNA
 XM_045421 Homo sapiens chromosome 20 open reading frame 194 (C20orf194), mRNA
 XM_045423 Homo sapiens KIAA0701 protein (KIAA0701), mRNA
 XM_045481 Homo sapiens likely ortholog of mouse 5-azacytidine induced gene 1 (AZI1),
 XM_045705 Homo sapiens similar to homologue of MJD, high homology to a genomic se
 XM_045712 Homo sapiens KIAA0316 gene product (KIAA0316), mRNA
 XM_045787 Homo sapiens hypothetical protein LOC92558 (LOC92558), mRNA
 XM_045792 Homo sapiens GCN1 general control of amino-acid synthesis 1-like 1 (yeast)
 XM_045907 Homo sapiens KIAA1170 protein (KIAA1170), mRNA
 XM_045911 Homo sapiens tomosyn-like (KIAA1006), mRNA
 XM_046099 Homo sapiens similar to small nuclear ribonucleoprotein E (LOC148064), mf
 XM_046284 Homo sapiens DKFZP434B172 protein (DKFZP434B172), mRNA
 XM_046305 Homo sapiens KIAA1205 (KIAA1205), mRNA
 XM_046390 Homo sapiens zinc finger protein 473 (ZNF473), mRNA
 XM_046437 Homo sapiens chromosome 20 open reading frame 50 (C20orf50), mRNA
 XM_046531 Homo sapiens KIAA1614 protein (KIAA1614), mRNA
 XM_046570 Homo sapiens KIAA1679 protein (KIAA1679), mRNA
 XM_046581 Homo sapiens zinc finger, SWIM domain containing 5 (ZSWIM5), mRNA
 XM_046800 Homo sapiens KIAA1272 protein (KIAA1272), mRNA
 XM_046877 Homo sapiens solute carrier family 39 (zinc transporter), member 14 (SLC39
 XM_046885 Homo sapiens KIAA1399 protein (KIAA1399), mRNA
 XM_046751 Homo sapiens protein tyrosine phosphatase, receptor type, f polypeptide (PT
 XM_046808 Homo sapiens neurofascin (NFASC), mRNA
 XM_046861 Homo sapiens KRAB box containing C2H2 type zinc finger bA526D8.4 (BA5;
 XM_047025 Homo sapiens ornithine aminotransferase-like 1 (OATL1), mRNA
 XM_047083 Homo sapiens similar to tubulin, beta 5 (LOC92755), mRNA
 XM_047214 Homo sapiens KIAA0930 protein (KIAA0930), mRNA
 XM_047325 Homo sapiens THO complex 2 (THOC2), mRNA
 XM_047355 Homo sapiens KIAA1765 protein (KIAA1765), mRNA
 XM_047357 Homo sapiens KIAA0342 gene product (KIAA0342), mRNA
 XM_047462 Homo sapiens Spir-2 protein (Spir-2), mRNA
 XM_047499 Homo sapiens hypothetical protein LOC149603 (LOC149603), mRNA
 XM_047550 Homo sapiens zinc finger protein 492 (ZNF492), mRNA
 XM_047554 Homo sapiens similar to Zinc finger protein 492 (LOC148198), mRNA
 XM_047610 Homo sapiens KIAA1086 (KIAA1086), mRNA

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XM_047617 Homo sapiens KIAA1349 protein (KIAA1349), mRNA
 XM_047707 Homo sapiens solute carrier family 39 (zinc transporter), member 10 (SLC39)
 XM_047734 Homo sapiens similar to hect domain and RLD 2 (LOC146489), mRNA
 XM_047770 Homo sapiens similar to alpha2-glucosyltransferase (LOC144245), mRNA
 XM_047995 Homo sapiens odd Oz/ten-m homolog 2 (ODZ2), mRNA
 XM_048070 Homo sapiens zinc finger protein 292 (ZNF292), mRNA
 XM_048104 Homo sapiens filaggrin (FLG), mRNA
 XM_048128 Homo sapiens KIAA1596 (KIAA1596), mRNA
 XM_048235 Homo sapiens Huntingtin interacting protein M (HYPM), mRNA
 XM_048362 Homo sapiens KIAA1543 (KIAA1543), mRNA
 XM_048462 Homo sapiens RUN and SH3 domain containing 2 (RUSC2), mRNA
 XM_048592 Homo sapiens KIAA1045 (KIAA1045), mRNA
 XM_048675 Homo sapiens KIAA1238 protein (KIAA1238), mRNA
 XM_048721 Homo sapiens hypothetical protein DKFPZp762K222 (DKFPZp762K222), mRNA
 XM_048747 Homo sapiens KIAA1223 protein (KIAA1223), mRNA
 XM_048774 Homo sapiens KIAA1332 protein (KIAA1332), mRNA
 XM_048786 Homo sapiens KIAA1061 protein (KIAA1061), mRNA
 XM_048825 Homo sapiens KIAA1026 protein (KIAA1026), mRNA
 XM_048898 Homo sapiens heat shock 70kDa protein 12A (HSPA12A), mRNA
 XM_049037 Homo sapiens trinucleotide repeat containing 9 (TNR09), mRNA
 XM_049078 Homo sapiens KIAA1239 protein (KIAA1239), mRNA
 XM_049237 Homo sapiens KIAA0841 (KIAA0841), mRNA
 XM_049349 Homo sapiens KIAA0534 protein (KIAA0534), mRNA
 XM_049351 Homo sapiens KIAA1600 protein (KIAA1600), mRNA
 XM_049380 Homo sapiens KIAA0339 gene product (KIAA0339), mRNA
 XM_049384 Homo sapiens chromosome 7 open reading frame 3 (C7orf3), mRNA
 XM_049575 Homo sapiens similar to succinate dehydrogenase flavoprotein subunit (LOC)
 XM_049619 Homo sapiens PR domain containing 6 (PRDM6), mRNA
 XM_049695 Homo sapiens vang-like 2 (van gogh, Drosophila) (VANGL2), mRNA
 XM_049952 Homo sapiens hypothetical protein FLJ23529 (FLJ23529), mRNA
 XM_050041 Homo sapiens myosin ID (MYO1D), mRNA
 XM_050219 Homo sapiens synaptopodin 2 (SYNPO2), mRNA
 XM_050278 Homo sapiens kinesin family member 26A (KIF26A), mRNA
 XM_050325 Homo sapiens KIAA1126 protein (KIAA1126), mRNA
 XM_050478 Homo sapiens KIAA1202 protein (KIAA1202), mRNA
 XM_050561 Homo sapiens SIN3 homolog B, transcriptional regulator (yeast) (SIN3B), mF
 XM_050564 Homo sapiens similar to RIKEN cDNA 2410004L22 gene (M. musculus) (MG)
 XM_050625 Homo sapiens secreted frizzled-related protein 2 (SFRP2), mRNA
 XM_050644 Homo sapiens KIAA1623 (KIAA1623), mRNA
 XM_050846 Homo sapiens Indian hedgehog homolog (Drosophila) (IHH), mRNA
 XM_051017 Homo sapiens KIAA0657 protein (KIAA0657), mRNA
 XM_051081 Homo sapiens KIAA0608 protein (KIAA0608), mRNA
 XM_051091 Homo sapiens KIAA1040 protein (KIAA1040), mRNA
 XM_051197 Homo sapiens KIAA1005 protein (KIAA1005), mRNA
 XM_051200 Homo sapiens fatso (FTO), mRNA
 XM_051221 Homo sapiens SPHK1 (sphingosine kinase type 1) interacting protein (SKIP)
 XM_051264 Homo sapiens thioredoxin reductase 3 (TXNRD3), mRNA
 XM_051271 Homo sapiens family with sequence similarity 10, member A6 (FAM10A6), m
 XM_051699 Homo sapiens KIAA1344 (KIAA1344), mRNA
 XM_051862 Homo sapiens hypothetical protein from EUROMIMAGE 588495 (LOC58489),
 XM_051956 Homo sapiens similar to KIAA0592 protein (LOC387680), mRNA
 XM_052561 Homo sapiens KIAA1337 protein (KIAA1337), mRNA
 XM_052597 Homo sapiens ubiquitin specific protease 53 (USP53), mRNA
 XM_052620 Homo sapiens mannosidase alpha class 2B member 2 (KIAA0935), mRNA
 XM_053074 Homo sapiens translocase of inner mitochondrial membrane 50 homolog (ye
 XM_053177 Homo sapiens similar to alpha tubulin (LOC112714), mRNA
 XM_053966 Homo sapiens hypothetical protein LOC113230 (LOC113230), mRNA

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XM_054284 Homo sapiens alpha-tubulin isotype H2-alpha (H2-ALPHA), mRNA
 XM_054313 Homo sapiens similar to T-complex protein 1 (LOC155100), mRNA
 XM_054983 Homo sapiens KIAA1952 protein (KIAA1952), mRNA
 XM_055095 Homo sapiens KIAA1906 protein (KIAA1906), mRNA
 XM_055481 Homo sapiens KIAA1915 protein (KIAA1915), mRNA
 XM_055636 Homo sapiens KIAA1912 protein (KIAA1912), mRNA
 XM_055725 Homo sapiens similar to unc-93 homolog B1; unc93 (C.elegans) homolog B;
 XM_055866 Homo sapiens lemur tyrosine kinase 3 (LMTK3), mRNA
 XM_056254 Homo sapiens heparan sulfate (glucosamine) 3-O-sulfotransferase 4 (HS3S1)
 XM_056282 Homo sapiens KIAA1904 protein (KIAA1904), mRNA
 XM_056298 Homo sapiens KIAA1889 protein (KIAA1889), mRNA
 XM_056434 Homo sapiens tetratricopeptide repeat domain 6 (TTC6), mRNA
 XM_056455 Homo sapiens Melanoma associated gene (D2S448), mRNA
 XM_056680 Homo sapiens hypothetical protein LOC115749 (LOC115749), mRNA
 XM_056681 Homo sapiens similar to ribosomal protein L14; 60S ribosomal protein L14 (L
 XM_057040 Homo sapiens KIAA1922 protein (KIAA1922), mRNA
 XM_057107 Homo sapiens KIAA1937 protein (KIAA1937), mRNA
 XM_057296 Homo sapiens hypothetical protein LOC116064 (LOC116064), mRNA
 XM_058332 Homo sapiens similar to hypothetical protein MGC45962 (LOC118670), mR
 XM_058335 Homo sapiens similar to ARF GTPase-activating protein (LOC118704), mR
 XM_058404 Homo sapiens hypothetical protein LOC119548 (LOC119548), mRNA
 XM_058426 Homo sapiens hypothetical protein FLJ00012 (FLJ00012), mRNA
 XM_058513 Homo sapiens hypothetical protein DKFPZ434H2111 (DKFPZ434H2111), mF
 XM_058581 Homo sapiens similar to hypothetical protein s530023G02 (LOC121642), mF
 XM_058611 Homo sapiens hypothetical LOC150928 (LOC150928), mRNA
 XM_058628 Homo sapiens chromosome 14 open reading frame 109 (C14orf109), mRNA
 XM_058661 Homo sapiens chromosome 14 open reading frame 35 (C14orf35), mRNA
 XM_058677 Homo sapiens similar to 60S ribosomal protein L21 (LOC123031), mRNA
 XM_058719 Homo sapiens similar to RIKEN cDNA C630028N24 gene (LOC123688), mR
 XM_058720 Homo sapiens similar to junction-mediating and regulatory protein p300 JMY
 XM_058721 Homo sapiens hypothetical protein LOC123722 (LOC123722), mRNA
 XM_058743 Homo sapiens hypothetical protein LOC123876 (LOC123876), mRNA
 XM_058857 Homo sapiens hypothetical LOC124871 (LOC124871), mRNA
 XM_058879 Homo sapiens hypothetical protein LOC124976 (LOC124976), mRNA
 XM_058931 Homo sapiens similar to hypothetical protein B230399E16 (LOC125704), mF
 XM_058956 Homo sapiens Purkinje cell protein 2 (PCP2), mRNA
 XM_058961 Homo sapiens trafficking protein particle complex 5 (TRAPPC5), mRNA
 XM_058964 Homo sapiens hypothetical protein LOC89887 (LOC89887), mRNA
 XM_058967 Homo sapiens similar to Elongation factor 1-delta (EF-1-delta) (Antigen NY-C
 XM_058987 Homo sapiens hypothetical protein LOC126167 (LOC126167), mRNA
 XM_058999 Homo sapiens hypothetical protein LOC126208 (LOC126208), mRNA
 XM_059037 Homo sapiens similar to LIM domains containing 1 (LOC126374), mRNA
 XM_059047 Homo sapiens hypothetical LOC126435 (LOC126435), mRNA
 XM_059051 Homo sapiens hypothetical protein LOC126520 (LOC126520), mRNA
 XM_059081 Homo sapiens hypothetical protein LOC126661 (LOC126661), mRNA
 XM_059074 Homo sapiens hypothetical protein LOC126755 (LOC126755), mRNA
 XM_059095 Homo sapiens formin binding protein 2 (FBNP2), mRNA
 XM_059104 Homo sapiens similar to CG5435-PA (LOC127003), mRNA
 XM_059132 Homo sapiens similar to RIKEN cDNA 4930549C01 (LOC127305), mRNA
 XM_059140 Homo sapiens similar to dJ39G22.2 (novel protein) (LOC127391), mRNA
 XM_059166 Homo sapiens similar to KIAA1697 protein (LOC127602), mRNA
 XM_059256 Homo sapiens hypothetical LOC128499 (LOC128499), mRNA
 XM_059267 Homo sapiens similar to RIKEN cDNA 2210009G21 (LOC128710), mRNA
 XM_059318 Homo sapiens KIAA1941 protein (KIAA1941), mRNA
 XM_059341 Homo sapiens hypothetical protein LOC129293 (LOC129293), mRNA
 XM_059368 Homo sapiens hypothetical protein LOC129607 (LOC129607), mRNA
 XM_059384 Homo sapiens hypothetical LOC129881 (LOC129881), mRNA

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XM_059396 Homo sapiens hypothetical LOC130063 (LOC130063), mRNA
 XM_059399 Homo sapiens similar to Calcium and integrin-binding protein 1 (Calmyrin) (C
 XM_059438 Homo sapiens similar to CG14894-PA (LOC130502), mRNA
 XM_059461 Homo sapiens similar to RIKEN cDNA A230078I05 gene (LOC130612), mR
 XM_059462 Homo sapiens hypothetical LOC130643 (LOC130643), mRNA
 XM_059473 Homo sapiens hypothetical LOC130839 (LOC130839), mRNA
 XM_059482 Homo sapiens FLJ00133 protein (FLJ00133), mRNA
 XM_059492 Homo sapiens hypothetical LOC131076 (LOC131076), mRNA
 XM_059548 Homo sapiens similar to SRSR846 (LOC131920), mRNA
 XM_059578 Homo sapiens similar to hypothetical protein A430083B19 (LOC132203), mF
 XM_059598 Homo sapiens similar to unc-93 homolog B1; unc93 (C.elegans) homolog B;
 XM_059608 Homo sapiens hypothetical LOC132870 (LOC132870), mRNA
 XM_059617 Homo sapiens similar to MGC69138 protein (LOC132946), mRNA
 XM_059669 Homo sapiens similar to ribosomal protein S3a; 40S ribosomal protein S3a;
 XM_059672 Homo sapiens hypothetical LOC133874 (LOC133874), mRNA
 XM_059689 Homo sapiens similar to 3110006E14Rik protein (LOC134111), mRNA
 XM_059702 Homo sapiens hypothetical protein FLJ36748 (FLJ36748), mRNA
 XM_059729 Homo sapiens interleukin-1 receptor-associated kinase 1 binding protein 1 (I
 XM_059730 Homo sapiens chromosome 6 open reading frame 159 (C6orf159), mRNA
 XM_059776 Homo sapiens FK506 binding protein 1C (FKBP1C), mRNA
 XM_059830 Homo sapiens similar to RIKEN cDNA 1700016G05 (LOC136242), mRNA
 XM_059832 Homo sapiens hypothetical protein LOC136288 (LOC136288), mRNA
 XM_059909 Homo sapiens hypothetical LOC137485 (LOC137485), mRNA
 XM_059923 Homo sapiens similar to Nuclear receptor binding factor-2 (LOC137829), mR
 XM_059929 Homo sapiens hypothetical protein LOC137886 (LOC137886), mRNA
 XM_059954 Homo sapiens chromosome 9 open reading frame 57 (C9orf57), mRNA
 XM_059956 Homo sapiens similar to RIKEN cDNA 1700028P14 (LOC138255), mRNA
 XM_059972 Homo sapiens hypothetical protein LOC138428 (LOC138428), mRNA
 XM_059987 Homo sapiens ankyrin repeat domain 19 (ANKRD19), mRNA
 XM_060020 Homo sapiens hypothetical protein BC016683 (LOC139231), mRNA
 XM_060054 Homo sapiens similar to XAGE-5 protein (LOC139793), mRNA
 XM_060087 Homo sapiens similar to template acylating factor-I alpha (LOC126598), mF
 XM_060104 Homo sapiens similar to RIKEN cDNA 543040H23 (LOC126637), mRNA
 XM_060171 Homo sapiens similar to erythrocyte membrane-associated giant protein anti
 XM_060278 Homo sapiens similar to tight junction protein 3 (zona occludens 3) (LOC126
 XM_060301 Homo sapiens similar to Olfactory receptor 2M6 (LOC127059), mRNA
 XM_060303 Homo sapiens similar to Olfactory receptor 2M6 (LOC127062), mRNA
 XM_060305 Homo sapiens similar to seven transmembrane helix receptor (LOC127064),
 XM_060307 Homo sapiens similar to Olfactory receptor 5BF1 (LOC127066), mRNA
 XM_060309 Homo sapiens similar to seven transmembrane helix receptor (LOC127068),
 XM_060310 Homo sapiens similar to seven transmembrane helix receptor (LOC127069),
 XM_060315 Homo sapiens similar to Olfactory receptor 2T4 (LOC127074), mRNA
 XM_060316 Homo sapiens olfactory receptor, family 2, subfamily T, member 1 (OR2T1),
 XM_060318 Homo sapiens similar to Olfactory receptor 2T11 (LOC127077), mRNA
 XM_060328 Homo sapiens similar to 60S ACIDIC RIBOSOMAL PROTEIN P1 (LOC12707
 XM_060417 Homo sapiens similar to 60S ribosomal protein L36 (LOC127295), mRNA
 XM_060458 Homo sapiens similar to Olfactory receptor 10J5 (LOC127385), mRNA
 XM_060509 Homo sapiens S100 calcium binding protein A7-like 2 (S100A7L2), mRNA
 XM_060535 Homo sapiens similar to ribosomal protein L18a; 60S ribosomal protein L18a
 XM_060537 Homo sapiens similar to IgB3 synthase (LOC127550), mRNA
 XM_060563 Homo sapiens similar to seven transmembrane helix receptor (LOC127608),
 XM_060569 Homo sapiens similar to seven transmembrane helix receptor (LOC127614),
 XM_060572 Homo sapiens similar to Olfactory receptor 5AV1 (LOC127617), mRNA
 XM_060580 Homo sapiens similar to olfactory receptor GA_x6K02SYHDF-1415-2371 (L
 XM_060597 Homo sapiens similar to zinc finger protein 135 (clone pHZ-17); zinc finger p
 XM_060880 Homo sapiens similar to dJ675G8.1 (novel zinc finger protein) (LOC128208),
 XM_060887 Homo sapiens similar to peptidyl-Pro cis trans isomerase (LOC128192), mR

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XM_060943 Homo sapiens similar to Nuclear transport factor 2 (NTF-2) (Placental protein
 XM_060945 Homo sapiens similar to Olfactory receptor 10T2 (LOC128360), mRNA
 XM_060951 Homo sapiens similar to Olfactory receptor 6P1 (LOC128366), mRNA
 XM_060952 Homo sapiens similar to seven transmembrane helix receptor (LOC128367),
 XM_060953 Homo sapiens similar to Olfactory receptor 10Z1 (LOC128368), mRNA
 XM_060955 Homo sapiens similar to seven transmembrane helix receptor (LOC128370),
 XM_060956 Homo sapiens similar to Olfactory receptor 6K6 (LOC128371), mRNA
 XM_060957 Homo sapiens similar to Olfactory receptor 6N1 (LOC128372), mRNA
 XM_060970 Homo sapiens paired related homeobox-like 1 (PRRXL1), mRNA
 XM_061055 Homo sapiens hypothetical protein FLJ32938 (FLJ32938), mRNA
 XM_061222 Homo sapiens similar to hypothetical protein 9930115F20 (LOC118934), mR
 XM_061427 Homo sapiens similar to Small nuclear ribonucleoprotein Sm D2 (snRNP con
 XM_061542 Homo sapiens similar to 40S ribosomal protein S8 (LOC119563), mRNA
 XM_061562 Homo sapiens similar to RIKEN cDNA 4632411J06 (LOC119593), mRNA
 XM_061610 Homo sapiens similar to Olfactory receptor 52E2 (LOC119678), mRNA
 XM_061611 Homo sapiens similar to Olfactory receptor 52J3 (LOC119679), mRNA
 XM_061614 Homo sapiens similar to Olfactory receptor 51L1 (LOC119682), mRNA
 XM_061619 Homo sapiens similar to Olfactory receptor 51A7 (LOC119687), mRNA
 XM_061624 Homo sapiens similar to Olfactory receptor 51S1 (LOC119692), mRNA
 XM_061626 Homo sapiens similar to Olfactory receptor 51F2 (LOC119694), mRNA
 XM_061627 Homo sapiens similar to Olfactory receptor 52R1 (LOC119695), mRNA
 XM_061628 Homo sapiens similar to seven transmembrane helix receptor (LOC119696),
 XM_061656 Homo sapiens similar to olfactory receptor MOR232-3 (LOC119749), mRNA
 XM_061666 Homo sapiens similar to Olfactory receptor 4X2 (LOC119764), mRNA
 XM_061674 Homo sapiens similar to seven transmembrane helix receptor (LOC119772),
 XM_061676 Homo sapiens similar to Olfactory receptor 52K2 (LOC119774), mRNA
 XM_061677 Homo sapiens similar to seven transmembrane helix receptor (LOC119775),
 XM_061849 Homo sapiens similar to FLJ10251 protein (LOC120082), mRNA
 XM_061884 Homo sapiens similar to fat3; fat3 protein (LOC120105), mRNA
 XM_061871 Homo sapiens FAT tumor suppressor homolog 3 (Drosophila) (FAT3), mRNA^F
 XM_061880 Homo sapiens similar to autoantigen NOR-90 (LOC120126), mRNA
 XM_061888 Homo sapiens similar to autoantigen NOR-90 (LOC120144), mRNA
 XM_061890 Homo sapiens similar to tripartite motif-containing 43 (LOC120146), mRNA
 XM_061930 Homo sapiens similar to Homeobox protein DBX1 (LOC120237), mRNA
 XM_062025 Homo sapiens similar to Hnrpa1 protein (LOC120364), mRNA
 XM_062162 Homo sapiens similar to Olfactory receptor 8I2 (LOC120586), mRNA
 XM_062263 Homo sapiens similar to seven transmembrane helix receptor (LOC120787),
 XM_062269 Homo sapiens similar to Olfactory receptor 56A4 (LOC120793), mRNA
 XM_062272 Homo sapiens similar to Olfactory receptor 56A1 (LOC120796), mRNA
 XM_062285 Homo sapiens similar to Olfactory receptor 2D3 (LOC120775), mRNA
 XM_062286 Homo sapiens similar to seven transmembrane helix receptor (LOC120776),
 XM_062300 Homo sapiens similar to RING finger protein 18 (Testis-specific ring-finger pr
 XM_062437 Homo sapiens similar to Keratin, type I cytoskeletal 18 (Cytokeratin 18) (K18
 XM_062467 Homo sapiens similar to seven transmembrane helix receptor (LOC121129),
 XM_062468 Homo sapiens similar to seven transmembrane helix receptor (LOC121130),
 XM_062520 Homo sapiens similar to Sucrase-isomaltase, intestinal (LOC121216), mRNA^F
 XM_062553 Homo sapiens similar to Olfactory receptor 10AD1 (LOC121275), mRNA
 XM_062594 Homo sapiens similar to seven transmembrane helix receptor (LOC121360),
 XM_062598 Homo sapiens similar to Olfactory receptor 10A7 (LOC121364), mRNA
 XM_062645 Homo sapiens similar to solute carrier family 9, member 7; nonselective sodi
 XM_062735 Homo sapiens forkhead box N4 (FOXN4), mRNA
 XM_062788 Homo sapiens similar to histidine-rich protein (LOC121792), mRNA
 XM_062871 Homo sapiens hypothetical protein FLJ40176 (FLJ40176), mRNA
 XM_062872 Homo sapiens hypothetical protein LOC121952 (LOC121952), mRNA
 XM_062890 Homo sapiens similar to peptidyl-Pro cis trans isomerase (LOC121981), mR
 XM_062912 Homo sapiens similar to Mitochondrial import receptor subunit TOM22 homol
 XM_062966 Homo sapiens similar to MAP/microtubule affinity-regulating kinase 3 (LOC1:

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XM_066176 Homo sapiens similar to bA218C14.1 (novel protein similar to mouse cystatrin)
XM_066177 Homo sapiens similar to bA218C14.1 (novel protein similar to mouse cystatrin)
XM_066189 Homo sapiens gamma-glutamyltransferase-like activity 3 (GGTLA3), mRNA
XM_066243 Homo sapiens hypothetical LOC128939 (LOC128939), mRNA
XM_066339 Homo sapiens hypothetical protein similar to topoisomerase (DNA) III beta (t-)
XM_066350 Homo sapiens similar to Ovis aries Y chromosome repeat region OY11.1 (3')
XM_066351 Homo sapiens hypothetical gene similar to gamma-glutamyltransferase-like 3
XM_066443 Homo sapiens similar to hypothetical protein MGC15827 (LOC139046), mRNA
XM_066452 Homo sapiens similar to plasmolipin (LOC139061), mRNA
XM_066457 Homo sapiens similar to SPANX N member 2 (LOC139067), mRNA
XM_066469 Homo sapiens similar to MAGE family testis and tumor-specific protein (LOC139116), mRNA
XM_066484 Homo sapiens similar to testis expressed sequence 13A (LOC139116), mRNA
XM_066534 Homo sapiens similar to Diacylglycerol kinase, delta (Diglyceride kinase) (DGK)
XM_066585 Homo sapiens similar to testis expressed sequence 13A (LOC139263), mRNA
XM_066621 Homo sapiens similar to envelope protein (LOC139302), mRNA
XM_066685 Homo sapiens similar to KIAA1387 protein (LOC139420), mRNA
XM_066690 Homo sapiens similar to H326 (LOC139425), mRNA
XM_066695 Homo sapiens similar to ferritin heavy chain - chicken (LOC139431), mRNA
XM_066701 Homo sapiens similar to melanoma antigen, family B, 4; melanoma-associated
XM_066752 Homo sapiens similar to E2F transcription factor 6 isoform a (LOC139542), mRNA
XM_066765 Homo sapiens similar to bA351K23.4 (novel protein) (LOC139562), mRNA
XM_066859 Homo sapiens similar to zinc finger protein 92 (LOC139735), mRNA
XM_066946 Homo sapiens hypothetical protein LOC139885 (LOC139886), mRNA
XM_067076 Homo sapiens similar to testis specific protein, Y-linked (LOC140103), mRNA
XM_067176 Homo sapiens similar to peptidyl-Pro cis trans isomerase (LOC131055), mRNA
XM_067193 Homo sapiens similar to microtubule-associated protein 6 (LOC131086), mRNA
XM_067228 Homo sapiens similar to otolin-1 (LOC131149), mRNA
XM_067369 Homo sapiens similar to abnormal cell LIN-41, heterochronic gene;
XM_067448 Homo sapiens similar to MEST (LOC131572), mRNA
XM_067503 Homo sapiens similar to peptidyl-Pro cis trans isomerase (LOC131691), mRNA
XM_067585 Homo sapiens hypothetical protein LOC131873 (LOC131873), mRNA
XM_067605 Homo sapiens similar to hypothetical protein (LOC131909), mRNA
XM_067904 Homo sapiens similar to Transcription factor BTF3 homolog 3 (LOC132556), mRNA
XM_067994 Homo sapiens similar to heat shock factor binding protein 1 (LOC132706), mRNA
XM_068121 Homo sapiens similar to homerin (LOC132969), mRNA
XM_068229 Homo sapiens similar to 953003A05 protein (LOC133185), mRNA
XM_068430 Homo sapiens similar to 60S acidic ribosomal protein P1 (LOC133609), mRNA
XM_068602 Homo sapiens hypothetical LOC133923 (LOC133923), mRNA
XM_068632 Homo sapiens similar to hypothetical protein MGC52498 (LOC133993), mRNA
XM_068681 Homo sapiens similar to seven transmembrane helix receptor (LOC134082), mRNA
XM_068682 Homo sapiens similar to Olfactory receptor 2Y1 (LOC134083), mRNA
XM_068889 Homo sapiens similar to eukaryotic translation initiation factor 3 subunit k; m
XM_068903 Homo sapiens similar to Ten-m2 (LOC134541), mRNA
XM_069035 Homo sapiens chromosome 6 open reading frame 213 (C6orf213), mRNA
XM_069595 Homo sapiens similar to Olfactory receptor 4F3 (LOC135896), mRNA
XM_069609 Homo sapiens similar to Olfactory receptor 9A2 (LOC135924), mRNA
XM_069612 Homo sapiens similar to OG-2 homeodomain protein-like; similar to U65067
XM_069616 Homo sapiens similar to seven transmembrane helix receptor (LOC135941), mRNA
XM_069619 Homo sapiens similar to olfactory receptor MOR261-13 (LOC135944), mRNA
XM_069621 Homo sapiens similar to Olfactory receptor 6B1 (Olfactory receptor 7-3) (OR7-3)
XM_069623 Homo sapiens similar to Olfactory receptor 2F2 (Olfactory receptor 7-1) (OR7-1)
XM_069728 Homo sapiens similar to beta-glucuronidase (LOC136132), mRNA
XM_069734 Homo sapiens similar to ribosomal protein L18; 60S ribosomal protein L18 (L18)
XM_069743 Homo sapiens similar to RIKEN cDNA 1500011L16 (LOC136157), mRNA
XM_069842 Homo sapiens similar to 60S ribosomal protein L15 (LOC136321), mRNA
XM_070233 Homo sapiens similar to ribosomal protein L10a (LOC137107), mRNA
XM_070277 Homo sapiens otoconin 90 (OC90), mRNA

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XM_070619 Homo sapiens similar to homeobox protein NKX2-6 (LOC137814), mRNA
XM_071013 Homo sapiens similar to bA62C3.1 (similar to testicular serine protease) (LO-
XM_071061 Homo sapiens AT rich interactive domain 3C (BRIGIT- like) (ARID3C), mRNA
XM_071093 Homo sapiens similar to Olfactory receptor 13C5 (LOC138799), mRNA
XM_071096 Homo sapiens similar to Olfactory receptor 13C8 (LOC138802), mRNA
XM_071097 Homo sapiens similar to Olfactory receptor 13C3 (LOC138803), mRNA
XM_071098 Homo sapiens similar to Olfactory receptor 13C4 (LOC138804), mRNA
XM_071099 Homo sapiens similar to Olfactory receptor 13F1 (LOC138805), mRNA
XM_071150 Homo sapiens similar to Olfactory receptor 1L8 (LOC138881), mRNA
XM_071151 Homo sapiens similar to Olfactory receptor 1N2 (LOC138882), mRNA
XM_071173 Homo sapiens similar to Hkr1p (LOC138932), mRNA
XM_071201 Homo sapiens similar to Centaurin gamma 2 (LOC138972), mRNA
XM_071712 Homo sapiens hypothetical protein LOC120376 (LOC120376), mRNA
XM_071793 Homo sapiens chromosome 14 open reading frame 28 (C14orf28), mRNA
XM_071866 Homo sapiens cerebellar degeneration-related protein 2, 62kDa (CDR2), mR
XM_072402 Homo sapiens aminoacylase 1-like 2 (ACY1L2), mRNA
XM_072554 Homo sapiens similar to RIKEN cDNA 4833436C18 gene (LOC138729), mR
XM_084000 Homo sapiens mitochondrial carrier triple repeat 2 (MCART2), mRNA
XM_084357 Homo sapiens similar to Hypothetical protein MGC56918 (LOC142827), mR
XM_084377 Homo sapiens similar to Triacylglycerol lipase, gastric precursor (Gastric lipa
XM_084445 Homo sapiens similar to ARF GTPase-activating protein (LOC143158), mRN
XM_084467 Homo sapiens similar to eukaryotic initiation factor 5A isoform I variant A (LC
XM_084482 Homo sapiens AT rich interactive domain 5B (MRF1-like) (ARID5B), mRNA
XM_084514 Homo sapiens heat shock 90kDa protein 1, alpha-like 3 (HSPCAL3), mRNA
XM_084529 Homo sapiens KIAA0298 gene product (KIAA0298), mRNA
XM_084530 Homo sapiens KIAA0033 protein (KIAA0033), mRNA
XM_084578 Homo sapiens PTPRF interacting protein, binding protein 2 (Iiprin beta 2) (PF
XM_084672 Homo sapiens similar to CDNA sequence BC021608 (LOC143941), mRNA
XM_084845 Homo sapiens similar to Interferon-induced transmembrane protein 3 (Interfe
XM_084852 Homo sapiens hypothetical LOC144404 (LOC144404), mRNA
XM_084868 Homo sapiens similar to MGC76214 protein (LOC144448), mRNA
XM_084990 Homo sapiens hypothetical LOC144962 (LOC144962), mRNA
XM_085028 Homo sapiens ATPase, Class VI, type 11A (ATP11A), mRNA
XM_085127 Homo sapiens KIAA0599 (KIAA0599), mRNA
XM_085138 Homo sapiens similar to ribosomal protein L3; 60S ribosomal protein L3; HIV
XM_085175 Homo sapiens tetratricopeptide repeat domain 7 like 1 (TTC7L1), mRNA
XM_085200 Homo sapiens hypothetical LOC145660 (LOC145660), mRNA
XM_085231 Homo sapiens hypothetical protein LOC145783 (LOC145783), mRNA
XM_085234 Homo sapiens unc-13 homolog C (C. elegans) (UNC13C), mRNA
XM_085236 Homo sapiens hypothetical LOC145788 (LOC145788), mRNA
XM_085261 Homo sapiens mesoderm posterior 2 (MESP2), mRNA
XM_085290 Homo sapiens similar to golgin-67 isoform c (LOC145988), mRNA
XM_085316 Homo sapiens similar to RIKEN cDNA 1810007E14; EST AA238765 (LOC14
XM_085347 Homo sapiens similar to hypothetical protein FLJ10815 (LOC146167), mRN/
XM_085367 Homo sapiens FLJ40162 protein (FLJ40162), mRNA
XM_085375 Homo sapiens zinc finger protein 90 homolog (mouse) (ZFP90), mRNA
XM_085383 Homo sapiens hypothetical protein LOC146206 (LOC146206), mRNA
XM_085403 Homo sapiens similar to CDNA sequence BC038613 (LOC146439), mRNA
XM_085507 Homo sapiens zinc finger protein 500 (ZNF500), mRNA
XM_085517 Homo sapiens hypothetical LOC146599 (LOC146599), mRNA
XM_085578 Homo sapiens FLJ46675 protein (FLJ46675), mRNA
XM_085596 Homo sapiens zinc finger protein 18 (KOX 11) (ZNF18), mRNA
XM_085606 Homo sapiens similar to CDRT15 protein (LOC146822), mRNA
XM_085634 Homo sapiens hypothetical protein LOC146909 (LOC146909), mRNA
XM_085689 Homo sapiens potassium channel tetramerisation domain containing 11 (KC
XM_085722 Homo sapiens similar to Tripartite motif protein 16 (Estrogen-responsive B pr
XM_085724 Homo sapiens hypothetical LOC147151 (LOC147151), mRNA

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XM_085775 Homo sapiens similar to 60S ribosomal protein L7a (Surfeit locus protein 3) (XM_085777
 Homo sapiens similar to additional sex combs like 2; polycomb group protein
 XM_085824 Homo sapiens hypothetical protein LOC147650 (LOC147650), mRNA
 XM_085830 Homo sapiens hypothetical protein LOC147649 (LOC147649), mRNA
 XM_085831 Homo sapiens hypothetical protein LOC147645 (LOC147645), mRNA
 XM_085833 Homo sapiens hypothetical protein LOC147646 (LOC147646), mRNA
 XM_085836 Homo sapiens KIAA1956 protein (KIAA1956), mRNA
 XM_085851 Homo sapiens similar to zinc finger protein 285 (LOC147711), mRNA
 XM_085870 Homo sapiens similar to complement C3 protein (GPC3) precursor (LOC147
 XM_085929 Homo sapiens Meis1, myeloid ecotropic viral integration site 1 homolog 3 (m
 XM_085932 Homo sapiens similar to VPRI645 (LOC147920), mRNA
 XM_085967 Homo sapiens hypothetical protein LOC147942 (LOC147942), mRNA
 XM_086001 Homo sapiens similar to Placental tissue protein 13 (Placenta protein 13) (G
 XM_086046 Homo sapiens hypothetical protein FLJ30663 (FLJ30663), mRNA
 XM_086095 Homo sapiens hypothetical protein LOC148203 (LOC148203), mRNA
 XM_086186 Homo sapiens hypothetical protein FLJ13815 (FLJ13815), mRNA
 XM_086188 Homo sapiens dnaj-like protein (LOC148418), mRNA
 XM_086257 Homo sapiens similar to ribosomal protein S15; rat insulinoma gene (LOC14:
 XM_086287 Homo sapiens similar to Osteotesticular phosphatase; protein tyrosine phosph
 XM_086308 Homo sapiens hypothetical protein LOC148766 (LOC148766), mRNA
 XM_086343 Homo sapiens similar to 60S ribosomal protein L17 (L23) (LOC148854), mR
 XM_086344 Homo sapiens similar to LIM homeo domain transcription factor (LOC148864
 XM_086360 Homo sapiens hypothetical protein LOC148915 (LOC148915), mRNA
 XM_086402 Homo sapiens hypothetical protein LOC149018 (LOC149018), mRNA
 XM_086409 Homo sapiens KIAA2025 protein (KIAA2025), mRNA
 XM_086494 Homo sapiens similar to similarity to monoubiquitin/carboxy-extension protei
 XM_086604 Homo sapiens similar to CHIA protein (LOC149620), mRNA
 XM_086616 Homo sapiens hypothetical protein LOC149643 (LOC149643), mRNA
 XM_086622 Homo sapiens hypothetical protein LOC149659 (LOC149659), mRNA
 XM_086637 Homo sapiens similar to RIKEN cDNA 1700049M11 (LOC149709), mRNA
 XM_086648 Homo sapiens similar to dJ579F20.1 (high-mobility group (nonhistone chrom
 XM_086650 Homo sapiens protein phosphatase 4, regulatory subunit 1-like (PPP4R1L), r
 XM_086725 Homo sapiens similar to bB329D4.2.1 (novel protein similar to a truncated nt
 XM_086732 Homo sapiens hypothetical protein LOC149950 (LOC149950), mRNA
 XM_086761 Homo sapiens hypothetical protein LOC150084 (LOC150084), mRNA
 XM_086828 Homo sapiens hypothetical protein LOC150368 (LOC150368), mRNA
 XM_086876 Homo sapiens similar to MGC5244 protein (LOC150207), mRNA
 XM_086879 Homo sapiens hypothetical protein LOC150371 (LOC150371), mRNA
 XM_086894 Homo sapiens hypothetical protein LOC150297 (LOC150297), mRNA
 XM_086905 Homo sapiens similar to RIKEN cDNA 2210021J22 (LOC150383), mRNA
 XM_086931 Homo sapiens similar to epsilon isoform of 14-3-3 protein (LOC150498), mR
 XM_086937 Homo sapiens similar to hypothetical protein A230046P18 (LOC150519), mF
 XM_086996 Homo sapiens hypothetical protein LOC150763 (LOC150763), mRNA
 XM_087056 Homo sapiens KIAA1841 protein (KIAA1841), mRNA
 XM_087062 Homo sapiens similar to 60S acidic ribosomal protein P1 (LOC150978), mR
 XM_087089 Homo sapiens KIAA0007 protein (KIAA0007), mRNA
 XM_087097 Homo sapiens hypothetical protein LOC151111 (LOC151111), mRNA
 XM_087137 Homo sapiens protein phosphatase 1 regulatory subunit 1A (LOC151242), m
 XM_087141 Homo sapiens hypothetical protein LOC151256 (LOC151256), mRNA
 XM_087167 Homo sapiens similar to KIAA1641 protein (LOC389008), mRNA
 XM_087171 Homo sapiens myeloid-associated differentiation marker-like (MYADML), mF
 XM_087182 Homo sapiens hypothetical protein LOC151363 (LOC151363), mRNA
 XM_087200 Homo sapiens hypothetical protein LOC151443 (LOC151443), mRNA
 XM_087208 Homo sapiens hypothetical protein LOC151451 (LOC151451), mRNA
 XM_087225 Homo sapiens similar to male-specific lethal 3-like 1 isoform a; drosophila M
 XM_087254 Homo sapiens ATPase, Class VI, type 11B (ATP11B), mRNA
 XM_087353 Homo sapiens KIAA0794 protein (KIAA0794), mRNA

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XM_087384 Homo sapiens hypothetical protein LOC152098 (LOC152098), mRNA
 XM_087386 Homo sapiens HEG homolog (HEG), mRNA
 XM_087483 Homo sapiens hypothetical protein LOC152519 (LOC152519), mRNA
 XM_087490 Homo sapiens similar to RIKEN cDNA 493343420 (LOC152586), mRNA
 XM_087499 Homo sapiens similar to 60S ribosomal protein L7a (Surfeit locus protein 3) (LOC152586)
 XM_087500 Homo sapiens similar to NEFA-interacting nuclear protein NIP30 (LOC15266)
 XM_087553 Homo sapiens similar to WW45 protein (LOC152891), mRNA
 XM_087593 Homo sapiens KIAA1430 protein (KIAA1430), mRNA
 XM_087671 Homo sapiens hypothetical LOC153441 (LOC153441), mRNA
 XM_087672 Homo sapiens KIAA1935 protein (KIAA1935), mRNA
 XM_087761 Homo sapiens similar to protein related with psoriasis (LOC153770), mRNA
 XM_087762 Homo sapiens hypothetical LOC153778 (LOC153778), mRNA
 XM_087800 Homo sapiens similar to CGI-62 protein (LOC153918), mRNA
 XM_087804 Homo sapiens synaptotagmin-like 3 (SYTL3), mRNA
 XM_087859 Homo sapiens similar to 60S ribosomal protein L21 (LOC154165), mRNA
 XM_087901 Homo sapiens similar to RIKEN cDNA 2410004A20 (LOC154288), mRNA
 XM_087928 Homo sapiens hypothetical protein LOC154449 (LOC154449), mRNA
 XM_088066 Homo sapiens similar to 60S ribosomal protein L35 (LOC154880), mRNA
 XM_088072 Homo sapiens hypothetical LOC154907 (LOC154907), mRNA
 XM_088118 Homo sapiens family with sequence similarity 10, member A7 (FAM10A7), m
 XM_088140 Homo sapiens hypothetical protein LOC155054 (LOC155054), mRNA
 XM_088142 Homo sapiens chromosome 7 open reading frame 32 (C7orf32), mRNA
 XM_088143 Homo sapiens similar to hypothetical protein 4931409K22 (LOC155046), mF
 XM_088315 Homo sapiens KIAA0870 protein (KIAA0870), mRNA
 XM_088331 Homo sapiens hypothetical protein LOC157570 (LOC157570), mRNA
 XM_088367 Homo sapiens similar to SPC18 protein (LOC157708), mRNA
 XM_088378 Homo sapiens chromosome 8 open reading frame 7 (C8orf7), mRNA
 XM_088459 Homo sapiens KIAA0310 (KIAA0310), mRNA
 XM_088491 Homo sapiens similar to Olfactory receptor 1Q1 (Olfactory receptor TPCR10
 XM_088516 Homo sapiens hypothetical LOC158226 (LOC158226), mRNA
 XM_088525 Homo sapiens chromosome 9 open reading frame 28 (C9orf28), mRNA
 XM_088551 Homo sapiens KIAA2026 (KIAA2026), mRNA
 XM_088566 Homo sapiens KIAA1958 (KIAA1958), mRNA
 XM_088567 Homo sapiens zinc finger protein 483 (ZNF483), mRNA
 XM_088578 Homo sapiens RAD26L hypothetical protein, alternatively spliced product; sIr
 XM_088636 Homo sapiens cyclin, basic protein of sperm head cytoskeleton 1 (CYLC1), i
 XM_088677 Homo sapiens similar to UPF3 regulator of nonsense transcripts homolog B i
 XM_088679 Homo sapiens hypothetical LOC158812 (LOC158812), mRNA
 XM_088680 Homo sapiens hypothetical LOC158813 (LOC158813), mRNA
 XM_088683 Homo sapiens similar to bA351K23.5 (novel protein) (LOC158835), mRNA
 XM_088684 Homo sapiens similar to Ab2-183 (LOC158830), mRNA
 XM_088686 Homo sapiens hypothetical LOC158825 (LOC158825), mRNA
 XM_088691 Homo sapiens hypothetical protein LOC158833 (LOC158833), mRNA
 XM_088726 Homo sapiens hypothetical LOC158957 (LOC158957), mRNA
 XM_088735 Homo sapiens hypothetical protein LOC158983 (LOC158983), mRNA
 XM_088768 Homo sapiens similar to F-box only protein 25 (LOC159176), mRNA
 XM_088797 Homo sapiens similar to BC035954 protein (LOC163301), mRNA
 XM_088817 Homo sapiens similar to Hypothetical protein DJ845O24.1 (LOC374949), mF
 XM_088951 Homo sapiens olfactomedin 3 (OLFM3), mRNA
 XM_089081 Homo sapiens deleted in neuroblastoma 5 (DNB5), mRNA
 XM_089243 Homo sapiens similar to cDNA sequence BC022623 (LOC163933), mRNA
 XM_089281 Homo sapiens similar to AD-003 protein (LOC149281), mRNA
 XM_089307 Homo sapiens similar to implantation-related RGS2-like protein (LOC164036
 XM_089384 Homo sapiens similar to RIKEN cDNA A430025D11 (LOC164118), mRNA
 XM_089747 Homo sapiens hypothetical protein FLJ35908 (FLJ35908), mRNA
 XM_089858 Homo sapiens similar to Olfactory receptor 52B4 (LOC143496), mRNA
 XM_089863 Homo sapiens similar to Olfactory receptor 5212 (LOC143502), mRNA

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XM_089866 Homo sapiens similar to RIKEN cDNA 1500011L16 (LOC143506), mRNA
 XM_090203 Homo sapiens similar to Olfactory receptor 2AG1 (HT3) (LOC144125), mRNA
 XM_090294 Homo sapiens hypothetical protein FLJ38508 (FLJ38508), mRNA
 XM_090844 Homo sapiens hypothetical protein LOC161291 (LOC161291), mRNA
 XM_090885 Homo sapiens chromosome 14 open reading frame 42 (C14orf42), mRNA
 XM_091156 Homo sapiens similar to Adenosine deaminase CG11994-PA (LOC161823),
 XM_091331 Homo sapiens hypothetical protein LOC162073 (LOC162073), mRNA
 XM_091809 Homo sapiens similar to WW domain binding protein 2 (LOC147468), mRNA
 XM_091830 Homo sapiens similar to G protein-coupled receptor 124 (LOC162835), mRNA
 XM_091886 Homo sapiens similar to Zinc finger protein Kr18 (HKr18) (LOC162962), mRNA
 XM_091914 Homo sapiens hypothetical protein LOC162993 (LOC162993), mRNA
 XM_092019 Homo sapiens similar to BC331191_1 (LOC163131), mRNA
 XM_092241 Homo sapiens similar to Olfactory receptor 8B3 (LOC150681), mRNA
 XM_092267 Homo sapiens similar to keratin 8; cytokeratin 8; keratin, type II cytoskeletal I
 XM_092342 Homo sapiens hypothetical protein FLJ39061 (FLJ39061), mRNA
 XM_092553 Homo sapiens similar to autoantigen NOR-90 (LOC151320), mRNA
 XM_092681 Homo sapiens similar to MLRQ subunit of the NADH ubiquinone oxidoreduct
 XM_092778 Homo sapiens hypothetical protein LOC164395 (LOC164395), mRNA
 XM_092995 Homo sapiens zinc finger protein 21 (KOX 14) (ZNF21), mRNA
 XM_093024 Homo sapiens hypothetical protein LOC169981 (LOC169981), mRNA
 XM_093087 Homo sapiens similar to transcription factor (p38 Interacting protein) (LOC17
 XM_093644 Homo sapiens similar to NACHT-, LRR- and PYD-containing protein 2 (PYRI
 XM_093813 Homo sapiens similar to hypothetical protein (LOC166348), mRNA
 XM_093839 Homo sapiens KIAA0826 protein (KIAA0826), mRNA
 XM_093895 Homo sapiens KIAA0882 protein (KIAA0882), mRNA
 XM_094066 Homo sapiens similar to RIKEN cDNA 5430419M09 (LOC152877), mRNA
 XM_094074 Homo sapiens similar to embryonic blastocoelar extracellular matrix protein p
 XM_094581 Homo sapiens SEC24 related gene family, member A (S. cerevisiae) (SEC24
 XM_094794 Homo sapiens dapper homolog 2, antagonist of beta-catenin (xenopus) (DAC
 XM_095568 Homo sapiens hypothetical protein DKFZp762C1112 (DKFZp762C1112), m
 XM_095746 Homo sapiens forkhead box D4 (FOXD4), mRNA
 XM_095965 Homo sapiens hypothetical protein LOC169834 (LOC169834), mRNA
 XM_095991 Homo sapiens chromosome 9 open reading frame 81 (C9orf81), mRNA
 XM_096317 Homo sapiens chromosome 10 open reading frame 73 (C10orf73), mRNA
 XM_096378 Homo sapiens hypothetical LOC143034 (LOC143034), mRNA
 XM_096472 Homo sapiens similar to RIKEN cDNA 1700021K07 (LOC143678), mRNA
 XM_096516 Homo sapiens hypothetical LOC143970 (LOC143970), mRNA
 XM_096642 Homo sapiens hypothetical LOC144631 (LOC144631), mRNA
 XM_096676 Homo sapiens hypothetical LOC144762 (LOC144762), mRNA
 XM_096688 Homo sapiens hypothetical protein LOC144920 (LOC144920), mRNA
 XM_096733 Homo sapiens chromosome 14 open reading frame 72 (C14orf72), mRNA
 XM_096734 Homo sapiens hypothetical LOC145197 (LOC145197), mRNA
 XM_096852 Homo sapiens hypothetical LOC145741 (LOC145741), mRNA
 XM_096864 Homo sapiens hypothetical LOC145780 (LOC145780), mRNA
 XM_096883 Homo sapiens hypothetical LOC145846 (LOC145846), mRNA
 XM_096885 Homo sapiens similar to ENSANGP00000021391 (LOC145853), mRNA
 XM_096919 Homo sapiens similar to SH2 domain protein 2A (T cell-specific adapter prot
 XM_097065 Homo sapiens hypothetical LOC146701 (LOC146701), mRNA
 XM_097265 Homo sapiens hypothetical protein LOC147670 (LOC147670), mRNA
 XM_097278 Homo sapiens hypothetical LOC147710 (LOC147710), mRNA
 XM_097347 Homo sapiens hypothetical LOC147941 (LOC147941), mRNA
 XM_097351 Homo sapiens hypothetical LOC147975 (LOC147975), mRNA
 XM_097580 Homo sapiens hypothetical protein LOC149086 (LOC149086), mRNA
 XM_097622 Homo sapiens similar to RIKEN cDNA C030014K22 gene (LOC149297), mR
 XM_097725 Homo sapiens hypothetical LOC149738 (LOC149738), mRNA
 XM_097729 Homo sapiens hypothetical LOC149704 (LOC149704), mRNA
 XM_097736 Homo sapiens chromosome 20 open reading frame 82 (C20orf82), mRNA

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XM_097753 Homo sapiens hypothetical LOC149913 (LOC149913), mRNA
 XM_097792 Homo sapiens hypothetical LOC150051 (LOC150051), mRNA
 XM_097886 Homo sapiens hypothetical protein LOC150223 (LOC150223), mRNA
 XM_097977 Homo sapiens hypothetical protein LOC150946 (LOC150946), mRNA
 XM_098008 Homo sapiens hypothetical LOC151154 (LOC151154), mRNA
 XM_098030 Homo sapiens hypothetical LOC151261 (LOC151261), mRNA
 XM_098117 Homo sapiens hypothetical LOC151760 (LOC151760), mRNA
 XM_098163 Homo sapiens hypothetical LOC152118 (LOC152118), mRNA
 XM_098164 Homo sapiens hypothetical LOC152122 (LOC152122), mRNA
 XM_098238 Homo sapiens SH3 domain protein D19 (DKFzp434D0215), mRNA
 XM_098317 Homo sapiens hypothetical LOC153134 (LOC153134), mRNA
 XM_098350 Homo sapiens hypothetical LOC153297 (LOC153297), mRNA
 XM_098368 Homo sapiens KIAA1317 protein (KIAA1317), mRNA
 XM_098406 Homo sapiens hypothetical LOC153630 (LOC153630), mRNA
 XM_098450 Homo sapiens hypothetical LOC153959 (LOC153959), mRNA
 XM_098512 Homo sapiens hypothetical LOC154323 (LOC154323), mRNA
 XM_098625 Homo sapiens hypothetical LOC154872 (LOC154872), mRNA
 XM_098762 Homo sapiens KIAA1416 protein (KIAA1416), mRNA
 XM_098828 Homo sapiens hypothetical LOC157813 (LOC157813), mRNA
 XM_098847 Homo sapiens hypothetical LOC157943 (LOC157943), mRNA
 XM_098940 Homo sapiens similar to zinc finger protein 11b (KOX 2) (LOC158431), mRNA
 XM_098980 Homo sapiens hypothetical LOC158730 (LOC158730), mRNA
 XM_099034 Homo sapiens hypothetical LOC159170 (LOC159170), mRNA
 XM_104657 Homo sapiens similar to RIKEN cDNA 1700019P01 (LOC164714), mRNA
 XM_106386 Homo sapiens KIAA1345 protein (KIAA1345), mRNA
 XM_113228 Homo sapiens similar to Proprotein convertase subtilisin/kexin type 7 precursor
 XM_113596 Homo sapiens similar to CG32542-PA (LOC196752), mRNA
 XM_113625 Homo sapiens hypothetical protein LOC195977 (LOC195977), mRNA
 XM_113641 Homo sapiens hypothetical protein LOC196051 (LOC196051), mRNA
 XM_113678 Homo sapiens nucleoporin 160kDa (NUP160), mRNA
 XM_113696 Homo sapiens hypothetical protein LOC196337 (LOC196337), mRNA
 XM_113706 Homo sapiens dynein, axonemal, heavy polypeptide 10 (DNAH10), mRNA
 XM_113708 Homo sapiens hypothetical protein LOC196394 (LOC196394), mRNA
 XM_113743 Homo sapiens hypothetical protein DKFzp313M0720 (DKFzp313M0720), mRNA
 XM_113763 Homo sapiens chromosome 14 open reading frame 125 (C14orf125), mRNA
 XM_113776 Homo sapiens hypothetical protein LOC196913 (LOC196913), mRNA
 XM_113796 Homo sapiens hypothetical protein LOC196996 (LOC196996), mRNA
 XM_113825 Homo sapiens similar to RIKEN cDNA 4930424G05 (LOC197135), mRNA
 XM_113871 Homo sapiens hypothetical protein LOC197350 (LOC197350), mRNA
 XM_113912 Homo sapiens similar to DKFZP568O084 protein (LOC201140), mRNA
 XM_113916 Homo sapiens similar to hypothetical protein A93006D11 (LOC201181), mRNA
 XM_113947 Homo sapiens KIAA0565 gene product (KIAA0565), mRNA
 XM_113967 Homo sapiens similar to Rab12 protein (LOC201475), mRNA
 XM_113971 Homo sapiens similar to B230208J24Rik protein (LOC201501), mRNA
 XM_113978 Homo sapiens hypothetical protein LOC284352 (LOC284352), mRNA
 XM_114000 Homo sapiens ankyrin repeat domain 24 (ANKRD24), mRNA
 XM_114047 Homo sapiens similar to KIAA0454 protein (LOC199882), mRNA
 XM_114067 Homo sapiens similar to expressed sequence AV028368 (LOC199953), mRNA
 XM_114087 Homo sapiens KIAA1836 protein (KIAA1836), mRNA
 XM_114090 Homo sapiens similar to KIAA0454 protein (LOC200019), mRNA
 XM_114129 Homo sapiens hypothetical LOC200159 (LOC200159), mRNA
 XM_114152 Homo sapiens hypothetical protein LOC200205 (LOC200205), mRNA
 XM_114156 Homo sapiens hypothetical protein LOC200213 (LOC200213), mRNA
 XM_114158 Homo sapiens similar to Polyadenylate-binding protein 2 (Poly(A)-binding protein)
 XM_114166 Homo sapiens similar to KIAA0386 (LOC200230), mRNA
 XM_114222 Homo sapiens similar to hypothetical protein (LOC200373), mRNA
 XM_114272 Homo sapiens selective LIM binding factor, rat homolog (SLB), mRNA

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XM_114297 Homo sapiens hypothetical protein FLJ10599 (FLJ10599), mRNA
 XM_114301 Homo sapiens similar to beta-1,4-mannosyltransferase; beta-1,4 mannosyltr
 XM_114303 Homo sapiens GRP1-binding protein GRSP1 (GRSP1), mRNA
 XM_114317 Homo sapiens similar to RIKEN cDNA 3110001N18 (LOC200916), mRNA
 XM_114355 Homo sapiens similar to esterase/N-deacetylase (EC 3.5.1.-), 50K hepatic - i
 XM_114415 Homo sapiens similar to Glycerol kinase, testis specific 1 (ATP:glycerol 3-ph
 XM_114418 Homo sapiens KIAA1729 protein (KIAA1729), mRNA
 XM_114430 Homo sapiens hypothetical protein LOC202051 (LOC202051), mRNA
 XM_114432 Homo sapiens KIAA1281 protein (KIAA1281), mRNA
 XM_114447 Homo sapiens KIAA1999 protein (KIAA1999), mRNA
 XM_114456 Homo sapiens hypothetical protein LOC202181 (LOC202181), mRNA
 XM_114481 Homo sapiens hypothetical LOC202404 (LOC202404), mRNA
 XM_114560 Homo sapiens similar to hypothetical protein MGC35361 (LOC202802), mRN
 XM_114611 Homo sapiens hypothetical protein KIAA1833 (KIAA1833), mRNA
 XM_114618 Homo sapiens hypothetical protein LOC203069 (LOC203069), mRNA
 XM_114621 Homo sapiens similar to RIKEN cDNA 4930578106 (LOC203076), mRNA
 XM_114685 Homo sapiens chromosome 9 open reading frame 21 (C9orf21), mRNA
 XM_114723 Homo sapiens similar to PAGE-5 protein (LOC203569), mRNA
 XM_114735 Homo sapiens complement component (3b/4b) receptor 1-like (CR1L), mRN
 XM_114973 Homo sapiens hypothetical protein LOC203806 (LOC203806), mRNA
 XM_114987 Homo sapiens similar to RIKEN cDNA 1500011L16 (LOC196120), mRNA
 XM_115009 Homo sapiens hypothetical protein LOC203859 (LOC203859), mRNA
 XM_115092 Homo sapiens similar to Olfactory receptor 56B4 (LOC196335), mRNA
 XM_115100 Homo sapiens similar to autoantigen NOR-90 (LOC196346), mRNA
 XM_115715 Homo sapiens similar to ENSANGP0000000189 (LOC200493), mRNA
 XM_115760 Homo sapiens similar to h2-calponin (LOC205272), mRNA
 XM_115789 Homo sapiens similar to chromosome 20 open reading frame 81 (LOC20059
 XM_115897 Homo sapiens similar to high mobility group protein homolog HMG4 (LOC200
 XM_115925 Homo sapiens similar to trophinin; melanoma antigen, family D, 3; trophinin-2
 XM_115974 Homo sapiens similar to hypothetical protein H41 (LOC200842), mRNA
 XM_116036 Homo sapiens similar to Gamma-aminobutyric-acid receptor rho-3 subunit pr
 XM_116384 Homo sapiens similar to Transcription initiation factor TFIID 28 kDa subunit (
 XM_116396 Homo sapiens similar to peptidylprolyl isomerase A (cyclophilin A) (LOC2022
 XM_116497 Homo sapiens chromosome 6 open reading frame 163 (C6orf163), mRNA
 XM_116623 Homo sapiens similar to Ubiquinol-cytochrome C reductase iron-sulfur subur
 XM_116936 Homo sapiens similar to RIKEN cDNA 2310038H17 (LOC196541), mRNA
 XM_116970 Homo sapiens similar to hypothetical protein (LOC196994), mRNA
 XM_116971 Homo sapiens hypothetical protein LOC196993 (LOC196993), mRNA
 XM_116980 Homo sapiens hypothetical LOC197049 (LOC197049), mRNA
 XM_117014 Homo sapiens hypothetical LOC197317 (LOC197317), mRNA
 XM_117030 Homo sapiens hypothetical LOC197387 (LOC197387), mRNA
 XM_117044 Homo sapiens hypothetical LOC201109 (LOC201109), mRNA
 XM_117056 Homo sapiens hypothetical LOC201201 (LOC201201), mRNA
 XM_117100 Homo sapiens hypothetical LOC201484 (LOC201484), mRNA
 XM_117112 Homo sapiens hypothetical LOC199680 (LOC199680), mRNA
 XM_117117 Homo sapiens hypothetical gene FLJ13072 (FLJ13072), mRNA
 XM_117152 Homo sapiens hypothetical LOC199897 (LOC199897), mRNA
 XM_117174 Homo sapiens hypothetical protein LOC200010 (LOC200010), mRNA
 XM_117213 Homo sapiens hypothetical LOC200292 (LOC200292), mRNA
 XM_117224 Homo sapiens similar to RIKEN cDNA 0610009J22 (LOC200312), mRNA
 XM_117236 Homo sapiens hypothetical LOC200475 (LOC200475), mRNA
 XM_117239 Homo sapiens hypothetical LOC200491 (LOC200491), mRNA
 XM_117257 Homo sapiens hypothetical LOC200624 (LOC200624), mRNA
 XM_117266 Homo sapiens hypothetical LOC200726 (LOC200726), mRNA
 XM_117268 Homo sapiens hypothetical LOC200731 (LOC200731), mRNA
 XM_117294 Homo sapiens hypothetical protein LOC200933 (LOC200933), mRNA
 XM_117408 Homo sapiens hypothetical LOC202546 (LOC202546), mRNA

XM_117451 Homo sapiens hypothetical LOC202775 (LOC202775), mRNA
 XM_117514 Homo sapiens hypothetical LOC203235 (LOC203235), mRNA
 XM_117548 Homo sapiens hypothetical LOC203413 (LOC203413), mRNA
 XM_165401 Homo sapiens similar to heat shock 70kD protein binding protein; progesterone
 XM_165448 Homo sapiens similar to BLOCK 23 (LOC220717), mRNA
 XM_165511 Homo sapiens similar to Fatty acid-binding protein, epidermal (E-FABP) (Psc
 XM_165534 Homo sapiens similar to methyltransferase-like protein 1 isoform c; D1075-II
 XM_165921 Homo sapiens similar to HP95 (LOC219392), mRNA
 XM_165973 Homo sapiens ubiquitin specific protease 24 (USP24), mRNA
 XM_166098 Homo sapiens neuregulin 3 (NRG3), mRNA
 XM_166090 Homo sapiens placenta-specific 9 (PLAC9), mRNA
 XM_166103 Homo sapiens DNA2 DNA replication helicase 2-like (yeast) (DNA2L), mRNA/
 XM_166110 Homo sapiens G protein-coupled receptor 158 (GPR158), mRNA
 XM_166125 Homo sapiens KIAA protein (similar to mouse paladin) (KIAA1274), mRNA
 XM_166132 Homo sapiens KIAA1462 (KIAA1462), mRNA
 XM_166138 Homo sapiens ankyrin repeat domain 16 (ANKRD16), mRNA
 XM_166140 Homo sapiens Scm-like with four mbt domains 2 (SFMBT2), mRNA
 XM_166160 Homo sapiens similar to chromosome 6 open reading frame 182 (LOC22101
 XM_166164 Homo sapiens hypothetical protein LOC219854 (LOC219854), mRNA
 XM_166203 Homo sapiens similar to RIKEN cDNA 1110030K22 (LOC219537), mRNA
 XM_166213 Homo sapiens KIAA0937 protein (KIAA0937), mRNA
 XM_166227 Homo sapiens macrophage expressed gene 1 (MPEG1), mRNA
 XM_166254 Homo sapiens odd Oz/ten-m homolog 4 (ODZ4), mRNA
 XM_166256 Homo sapiens microtubule-associated protein 6 (MAP6), mRNA
 XM_166270 Homo sapiens KIAA0774 (KIAA0774), mRNA
 XM_166300 Homo sapiens absent in melanoma 1 (AIM1), mRNA
 XM_166320 Homo sapiens KIAA1553 (KIAA1553), mRNA
 XM_166346 Homo sapiens chromosome 6 open reading frame 129 (C6orf129), mRNA
 XM_166372 Homo sapiens leucine rich repeat and fibronectin type III domain containing :
 XM_166376 Homo sapiens KIAA1949 protein (KIAA1949), mRNA
 XM_166420 Homo sapiens RPEL repeat containing 1 (RPEL1), mRNA
 XM_166432 Homo sapiens hypothetical protein LOC221442 (LOC221442), mRNA
 XM_166443 Homo sapiens tudor domain containing 6 (TDRD6), mRNA
 XM_166450 Homo sapiens bromodomain and PHD finger containing, 3 (BRPF3), mRNA
 XM_166451 Homo sapiens KIAA1588 (KIAA1588), mRNA
 XM_166453 Homo sapiens tau tubulin kinase 1 (TTBK1), mRNA
 XM_166479 Homo sapiens KIAA0240 (KIAA0240), mRNA
 XM_166508 Homo sapiens TWIST neighbor (TWISTNB), mRNA
 XM_166523 Homo sapiens tweety homolog 3 (Drosophila) (TTYH3), mRNA
 XM_166527 Homo sapiens KIAA0415 gene product (KIAA0415), mRNA
 XM_166529 Homo sapiens glucocorticoid induced transcript 1 (GLCCH1), mRNA
 XM_166532 Homo sapiens KIAA1950 protein (KIAA1950), mRNA
 XM_166571 Homo sapiens KIAA0363 protein (KIAA0363), mRNA
 XM_166573 Homo sapiens KIAA0895 protein (KIAA0895), mRNA
 XM_166630 Homo sapiens similar to KIAA2020 protein (LOC387692), mRNA
 XM_166659 Homo sapiens hypothetical protein LOC220213 (LOC220213), mRNA
 XM_166707 Homo sapiens olfactory receptor, family 13, subfamily A, member 1 (OR13A
 XM_166720 Homo sapiens similar to Protein C21orf58 (LOC220988), mRNA
 XM_166747 Homo sapiens similar to KIAA1838 (LOC219797), mRNA
 XM_166757 Homo sapiens similar to Olfactory receptor 8B12 (LOC219858), mRNA
 XM_166787 Homo sapiens similar to seven transmembrane helix receptor (LOC219865),
 XM_166776 Homo sapiens similar to Olfactory receptor 10G8 (LOC219869), mRNA
 XM_166777 Homo sapiens similar to Olfactory receptor 10G9 (LOC219870), mRNA
 XM_166780 Homo sapiens similar to Olfactory receptor 10S1 (LOC219873), mRNA
 XM_166781 Homo sapiens similar to Olfactory receptor 6T1 (LOC219874), mRNA
 XM_166782 Homo sapiens similar to Olfactory receptor 4D5 (LOC219875), mRNA
 XM_166805 Homo sapiens similar to seven transmembrane helix receptor (LOC219417),

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XM_166808 Homo sapiens similar to Olfactory receptor 5A51 (LOC219447), mRNA
 XM_166813 Homo sapiens similar to Olfactory receptor 8K5 (LOC219453), mRNA
 XM_166818 Homo sapiens similar to hypothetical protein FLJ13194 (LOC219462), mRNA
 XM_166820 Homo sapiens similar to Olfactory receptor 5T2 (LOC219464), mRNA
 XM_166823 Homo sapiens similar to Olfactory receptor 8H1 (LOC219469), mRNA
 XM_166825 Homo sapiens similar to Olfactory receptor 8K3 (LOC219473), mRNA
 XM_166829 Homo sapiens similar to Olfactory receptor 8J1 (LOC219477), mRNA
 XM_166831 Homo sapiens similar to seven transmembrane helix receptor (LOC219479), mRNA
 XM_166834 Homo sapiens similar to Olfactory receptor 5M3 (LOC219482), mRNA
 XM_166835 Homo sapiens similar to Olfactory receptor 5M8 (LOC219484), mRNA
 XM_166845 Homo sapiens similar to Olfactory receptor 5AR1 (LOC219493), mRNA
 XM_166856 Homo sapiens similar to expressed sequence AB41794 (LOC219527), mRNA
 XM_166888 Homo sapiens similar to Olfactory receptor 4C16 (LOC219428), mRNA
 XM_166869 Homo sapiens similar to seven transmembrane helix receptor (LOC219429), mRNA
 XM_166871 Homo sapiens similar to Olfactory receptor 4S2 (LOC219431), mRNA
 XM_166872 Homo sapiens similar to Olfactory receptor 4C8 (LOC219432), mRNA
 XM_166877 Homo sapiens similar to Olfactory receptor 5D14 (LOC219436), mRNA
 XM_166878 Homo sapiens similar to Olfactory receptor 5L1 (OST262) (LOC219437), mRNA
 XM_166879 Homo sapiens similar to Olfactory receptor 5D18 (LOC219438), mRNA
 XM_166898 Homo sapiens similar to Olfactory receptor 5A2 (LOC219981), mRNA
 XM_166899 Homo sapiens similar to Olfactory receptor 5A1 (OST181) (LOC219982), mRNA
 XM_166900 Homo sapiens similar to Olfactory receptor 4D6 (LOC219983), mRNA
 XM_166903 Homo sapiens similar to seven transmembrane helix receptor (LOC219986), mRNA
 XM_166910 Homo sapiens similar to Olfactory receptor 6Q1 (LOC219952), mRNA
 XM_166912 Homo sapiens similar to Olfactory receptor 9I1 (LOC219954), mRNA
 XM_166914 Homo sapiens similar to Olfactory receptor 9Q1 (LOC219956), mRNA
 XM_166915 Homo sapiens similar to seven transmembrane helix receptor (LOC219957), mRNA
 XM_166916 Homo sapiens similar to Olfactory receptor 1S2 (LOC219958), mRNA
 XM_166917 Homo sapiens similar to Olfactory receptor 1S1 (LOC219959), mRNA
 XM_166918 Homo sapiens similar to Olfactory receptor 10Q1 (LOC219960), mRNA
 XM_166923 Homo sapiens similar to Olfactory receptor 5B17 (LOC219965), mRNA
 XM_166926 Homo sapiens similar to olfactory receptor MOR202-4 (LOC219968), mRNA
 XM_166986 Homo sapiens similar to Meningioma-expressed antigen 6/11 (MEA6) (MEA1
 XM_166971 Homo sapiens similar to Leucine-rich repeat protein SHOC-2 (Ras-binding p
 XM_167001 Homo sapiens similar to 40S ribosomal protein S26 (LOC219542), mRNA
 XM_167044 Homo sapiens solute carrier family 35, member F1 (SLC35F1), mRNA
 XM_167072 Homo sapiens benzodiazepine receptor (peripheral)-like 1 (BZRP1), mRNA
 XM_167147 Homo sapiens zinc finger protein 390 (ZNF390), mRNA
 XM_167149 Homo sapiens chromosome 6 open reading frame 194 (C6orf194), mRNA
 XM_167152 Homo sapiens similar to Zinc finger protein 192 (LD5-1) (LOC222701), mRNA
 XM_167254 Homo sapiens similar to tropomyosin 3 (LOC221875), mRNA
 XM_167275 Homo sapiens similar to ribosomal protein L23 (LOC222901), mRNA
 XM_167709 Homo sapiens hypothetical protein LOC221061 (LOC221061), mRNA
 XM_167711 Homo sapiens integrin, alpha 8 (ITGA8), mRNA
 XM_167908 Homo sapiens hypothetical LOC221140 (LOC221140), mRNA
 XM_168030 Homo sapiens zinc finger protein 319 (ZNF319), mRNA
 XM_168053 Homo sapiens chromosome 6 open reading frame 184 (C6orf184), mRNA
 XM_168055 Homo sapiens chromosome 6 open reading frame 185 (C6orf185), mRNA
 XM_168080 Homo sapiens chromosome 6 open reading frame 154 (C6orf154), mRNA
 XM_168073 Homo sapiens hypothetical LOC221344 (LOC221344), mRNA
 XM_168101 Homo sapiens hypothetical LOC221766 (LOC221766), mRNA
 XM_168223 Homo sapiens similar to RIKEN cDNA A530018O06 gene (LOC221813), mRNA
 XM_168302 Homo sapiens zinc finger protein 36 (KOX 18) (ZNF36), mRNA
 XM_168354 Homo sapiens similar to SMT3 suppressor of mif two 3 homolog 2 (LOC222C
 XM_168521 Homo sapiens hypothetical LOC222190 (LOC222190), mRNA
 XM_168576 Homo sapiens mucin 3B (MUC3B), mRNA
 XM_168583 Homo sapiens mucin 17 (MUC17), mRNA

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XM_168585 Homo sapiens similar to mucin 11 (LOC219612), mRNA
 XM_168590 Homo sapiens zotofin related factor 1 (ZRF1), mRNA
 XM_169057 Homo sapiens hypothetical LOC219908 (LOC219908), mRNA
 XM_169227 Homo sapiens similar to proline rich antigen 2 (LOC220061), mRNA
 XM_169258 Homo sapiens similar to KIAA0543 protein (LOC219638), mRNA
 XM_169434 Homo sapiens similar to TAR DNA binding protein (LOC219414), mRNA
 XM_170525 Homo sapiens similar to ARF GTPase-activating protein (LOC255319), mRNA
 XM_170536 Homo sapiens similar to eukaryotic translation initiation factor 2, subunit 3 gα
 XM_170597 Homo sapiens similar to peptidylprolyl isomerase A (LOC256374), mRNA
 XM_170620 Homo sapiens similar to 60S ribosomal protein L21 (LOC256457), mRNA
 XM_170637 Homo sapiens similar to beta-tubulin 4Q (LOC253936), mRNA
 XM_170658 Homo sapiens tangerin (DKFZp762C186), mRNA
 XM_170659 Homo sapiens similar to 2010003K11Rik protein (LOC254439), mRNA
 XM_170667 Homo sapiens hypothetical protein LOC254359 (LOC254359), mRNA
 XM_170708 Homo sapiens hypothetical LOC255411 (LOC255411), mRNA
 XM_170736 Homo sapiens hypothetical protein LOC255312 (LOC255312), mRNA
 XM_170749 Homo sapiens similar to GARNL1 protein (LOC387984), mRNA
 XM_170754 Homo sapiens similar to serine (or cysteine) proteinase inhibitor, clade A (α1)
 XM_170777 Homo sapiens similar to Microsomal signal peptidase 25 kDa subunit (SPase
 XM_170840 Homo sapiens similar to CDRT15 protein (LOC256223), mRNA
 XM_170842 Homo sapiens hypothetical protein FLJ40244 (FLJ40244), mRNA
 XM_170909 Homo sapiens similar to hypothetical protein MGC20470 (LOC257177), mRNA
 XM_170950 Homo sapiens similar to OSJNBa0043A12.32 (LOC254897), mRNA
 XM_170994 Homo sapiens hypothetical LOC255349 (LOC255349), mRNA
 XM_171008 Homo sapiens similar to high mobility group AT-hook 1 (LOC257200), mRNA
 XM_171013 Homo sapiens similar to Gamma-2-syntrophin (G2SYN) (Syntrophin 5) (SYN
 XM_171032 Homo sapiens hypothetical protein LOC255812 (LOC255812), mRNA
 XM_171040 Homo sapiens similar to MEGF6 (LOC253820), mRNA
 XM_171054 Homo sapiens KIAA0527 protein (KIAA0527), mRNA
 XM_171080 Homo sapiens hypothetical protein MGC50836 (MGC50836), mRNA
 XM_171068 Homo sapiens hypothetical protein LOC253017 (LOC253017), mRNA
 XM_171078 Homo sapiens similar to ALGV3072 (LOC255324), mRNA
 XM_171081 Homo sapiens similar to ras-related C3 botulinum toxin substrate 1 Isoform F
 XM_171094 Homo sapiens similar to eukaryotic translation initiation factor eIF4E-1 (LOC
 XM_171105 Homo sapiens hypothetical LOC255338 (LOC255338), mRNA
 XM_171150 Homo sapiens similar to bA145E8.1 (KIAA1074) (LOC254027), mRNA
 XM_171154 Homo sapiens similar to myelin protein zero-like 1; protein zero related (LOC
 XM_171156 Homo sapiens similar to bA145E8.1 (KIAA1074) (LOC254750), mRNA
 XM_171158 Homo sapiens similar to ribosomal protein S2; 40S ribosomal protein S2 (MC
 XM_171183 Homo sapiens similar to Six transmembrane epithelial antigen of prostate (L
 XM_171165 Homo sapiens similar to cAMP-dependent protein kinase type I-beta regulat
 XM_171171 Homo sapiens similar to hypothetical protein MGC49416 (LOC255374), mRNA
 XM_171207 Homo sapiens similar to RIKEN cDNA 4930429J24 (LOC255220), mRNA
 XM_171224 Homo sapiens coactivator associated arginine methyltransferase 1-like (CA
 XM_171410 Homo sapiens hypothetical protein DKFZp667B0210 (DKFZp667B0210), m
 XM_171424 Homo sapiens similar to olfactory receptor (LOC256892), mRNA
 XM_171447 Homo sapiens similar to Transcription elongation factor B polypeptide 2 (RN
 XM_171489 Homo sapiens similar to seven transmembrane helix receptor (LOC256144),
 XM_171480 Homo sapiens similar to Olfactory receptor 4S1 (LOC256148), mRNA
 XM_171485 Homo sapiens similar to Claudin-22 (LOC255244), mRNA
 XM_171503 Homo sapiens similar to large subunit ribosomal protein L36a (LOC255701),
 XM_171528 Homo sapiens similar to seven transmembrane helix receptor (LOC255725),
 XM_171536 Homo sapiens mas-related G protein-coupled MRGE (MRGE), mRNA
 XM_171578 Homo sapiens similar to olfactory receptor MOR109-1 (LOC254783), mRNA
 XM_171581 Homo sapiens similar to olfactory receptor MOR111-4 (LOC254786), mRNA
 XM_171590 Homo sapiens similar to ribosomal protein L22 (LOC256441), mRNA
 XM_171766 Homo sapiens similar to high mobility group protein homolog HMG4 (LOC25

XM_171855 Homo sapiens similar to myeloid-associated differentiation marker (LOC2552
 XM_171892 Homo sapiens similar to ribosomal protein L31 (LOC253013), mRNA
 XM_171973 Homo sapiens similar to zinc finger protein 91 (HPF7, HTF10) (LOC253342),
 XM_172230 Homo sapiens similar to ribosomal protein S17 (LOC257039), mRNA
 XM_172341 Homo sapiens hypothetical protein FLJ35036 (FLJ35036), mRNA
 XM_172389 Homo sapiens similar to Tryptophan-rich protein (Congenital heart disease 5
 XM_172751 Homo sapiens similar to Olfactory receptor 1L4 (Olfactory receptor 9-E) (OR1
 XM_172801 Homo sapiens KIAA1210 protein (KIAA1210), mRNA
 XM_172852 Homo sapiens hypothetical LOC256676 (LOC256676), mRNA
 XM_172855 Homo sapiens hypothetical LOC255849 (LOC255849), mRNA
 XM_172860 Homo sapiens hypothetical LOC255649 (LOC255649), mRNA
 XM_172861 Homo sapiens similar to ZP1 precursor (LOC255714), mRNA
 XM_172874 Homo sapiens hypothetical LOC253724 (LOC253724), mRNA
 XM_172889 Homo sapiens hypothetical LOC256176 (LOC256176), mRNA
 XM_172917 Homo sapiens hypothetical LOC256453 (LOC256453), mRNA
 XM_172929 Homo sapiens hypothetical protein LOC255189 (LOC255189), mRNA
 XM_172931 Homo sapiens hypothetical protein LOC254559 (LOC254559), mRNA
 XM_172968 Homo sapiens hypothetical protein LOC253962 (LOC253962), mRNA
 XM_172995 Homo sapiens hypothetical LOC255809 (LOC255809), mRNA
 XM_173012 Homo sapiens hypothetical LOC256686 (LOC256686), mRNA
 XM_173015 Homo sapiens hypothetical LOC256483 (LOC256483), mRNA
 XM_173036 Homo sapiens hypothetical protein LOC255654 (LOC255654), mRNA
 XM_173063 Homo sapiens hypothetical LOC253662 (LOC253662), mRNA
 XM_173068 Homo sapiens hypothetical LOC253584 (LOC253584), mRNA
 XM_173083 Homo sapiens hypothetical protein LOC255025 (LOC255025), mRNA
 XM_173084 Homo sapiens hypothetical protein LOC254827 (LOC254827), mRNA
 XM_173087 Homo sapiens hypothetical protein LOC255798 (LOC255798), mRNA
 XM_173105 Homo sapiens hypothetical LOC256283 (LOC256283), mRNA
 XM_173119 Homo sapiens hypothetical LOC255130 (LOC255130), mRNA
 XM_173120 Homo sapiens hypothetical LOC254938 (LOC254938), mRNA
 XM_173132 Homo sapiens similar to unc-93 homolog B1; unc93 (C.elegans) homolog B;
 XM_173135 Homo sapiens hypothetical LOC256880 (LOC256880), mRNA
 XM_173140 Homo sapiens hypothetical LOC253254 (LOC253254), mRNA
 XM_173160 Homo sapiens hypothetical LOC255187 (LOC255187), mRNA
 XM_173164 Homo sapiens hypothetical LOC256096 (LOC256096), mRNA
 XM_173166 Homo sapiens chromosome 6 open reading frame 191 (C6orf191), mRNA
 XM_173173 Homo sapiens amine oxidase, flavin containing 1 (AOX1), mRNA
 XM_175125 Homo sapiens hemiceritin-2 (DKFPZp434P0216), mRNA
 XM_208028 Homo sapiens similar to double homeobox protein (LOC283039), mRNA
 XM_208035 Homo sapiens similar to hypothetical protein FLJ10817 (LOC347806), mRNA
 XM_208042 Homo sapiens similar to 40S ribosomal protein S25 (LOC283114), mRNA
 XM_208043 Homo sapiens similar to RING finger protein 18 (Testis-specific ring-finger pr
 XM_208058 Homo sapiens similar to NADPH oxidase 4 variant (LOC283247), mRNA
 XM_208060 Homo sapiens similar to tripartite motif-containing 51 (LOC283257), mRNA
 XM_208061 Homo sapiens similar to tripartite motif protein 48; TRIM48 (LOC283259), m
 XM_208072 Homo sapiens similar to ribosomal protein L13a; 60S ribosomal protein L13a
 XM_208080 Homo sapiens similar to Polyhomeotic-like protein 1 (Early development regu
 XM_208097 Homo sapiens similar to telomeric repeat binding factor 1 isoform 2: Telomer
 XM_208108 Homo sapiens similar to synaptogyrin 2; cellugyrin (LOC283698), mRNA
 XM_208116 Homo sapiens similar to elongation factor SIII p15 subunit (LOC283747), mR
 XM_208142 Homo sapiens similar to hypothetical protein (LOC283957), mRNA
 XM_208145 Homo sapiens similar to rhophilin-like protein; RhoB effector; rhophilin-2; rho
 XM_208151 Homo sapiens similar to solute carrier family 16, member 6; monocarboxylate
 XM_208185 Homo sapiens similar to large subunit ribosomal protein L36a (LOC284230),
 XM_208200 Homo sapiens similar to Heterogeneous nuclear ribonucleoprotein A1 (Helix-
 XM_208203 Homo sapiens similar to methyl-CpG binding domain protein 3-like 2 (LOC28
 XM_208204 Homo sapiens similar to actin-related protein 2 (LOC284441), mRNA

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XM_208213 Homo sapiens similar to fatty acid omega-hydroxylase CYP4A11 (LOC28454)
 XM_208234 Homo sapiens similar to inactive progesterone receptor, 23 kD; likely orthok
 XM_208250 Homo sapiens similar to FRG1 protein (FSHD region gene 1 protein) (LOC28
 XM_208261 Homo sapiens similar to lymphocyte activation-associated protein; kelch (Drc
 XM_208270 Homo sapiens acrosin-like protease (ba395L14.13), mRNA
 XM_208281 Homo sapiens similar to ribosomal protein L18a; 60S ribosomal protein L18a
 XM_208300 Homo sapiens similar to 60S ribosomal protein L23a (LOC285214), mRNA
 XM_208312 Homo sapiens similar to unc-93 homolog B1; unc93 (C.elegans) homolog B;
 XM_208313 Homo sapiens similar to tropomyosin 4 (LOC285321), mRNA
 XM_208316 Homo sapiens similar to vascular endothelial zinc finger 1 (LOC285388), mR
 XM_208317 Homo sapiens similar to 40S ribosomal protein S20 (LOC285384), mRNA
 XM_208319 Homo sapiens similar to Epidermal Langerhans cell protein LCP1 (LOC2854
 XM_208320 Homo sapiens similar to mitochondrial translational release factor 1-like (LOC
 XM_208333 Homo sapiens hypothetical protein MGC48637 (MGC48637), mRNA
 XM_208352 Homo sapiens alcohol dehydrogenase 5B (ADH5B), mRNA
 XM_208356 Homo sapiens similar to cytochrome c oxidase subunit VIa polypeptide 1 pre
 XM_208361 Homo sapiens similar to RPL6 protein (LOC285900), mRNA
 XM_208373 Homo sapiens similar to Heterogeneous nuclear ribonucleoprotein A1 (Helix-
 XM_208403 Homo sapiens similar to Von Ebners gland protein precursor (VEG protein) (C
 XM_208423 Homo sapiens similar to ribosomal protein S2; 40S ribosomal protein S2 (LO
 XM_208431 Homo sapiens similar to ubiquitin-conjugating enzyme UbcM2 (LOC286480),
 XM_208438 Homo sapiens similar to Tetratricopeptide repeat protein 3 (TPR repeat prote
 XM_208443 Homo sapiens Ras-like GTPase-like (LOC286526), mRNA
 XM_208502 Homo sapiens similar to hypothetical protein 9330140G23 (LOC283071), mF
 XM_208522 Homo sapiens KIAA1394 protein (KIAA1394), mRNA
 XM_208524 Homo sapiens hypothetical protein LOC283129 (LOC283129), mRNA
 XM_208529 Homo sapiens hypothetical protein DKFZp547C195 (DKFZp547C195), mRN
 XM_208541 Homo sapiens similar to Olfactory receptor 8D2 (Olfactory receptor-like prote
 XM_208543 Homo sapiens similar to Olfactory receptor 8D1 (Olfactory receptor-like prote
 XM_208545 Homo sapiens similar to bone morphogenetic protein receptor, type IA precu
 XM_208554 Homo sapiens similar to Protein farnesyltransferase/geranylgeranyltransferase
 XM_208563 Homo sapiens hypothetical LOC283202 (LOC283202), mRNA
 XM_208604 Homo sapiens similar to Olfactory receptor 10A4 (HP2) (Olfactory receptor-III
 XM_208612 Homo sapiens hypothetical LOC283321 (LOC283321), mRNA
 XM_208613 Homo sapiens similar to 60 kDa heat shock protein, mitochondrial precursor
 XM_208635 Homo sapiens similar to cDNA sequence BC030307 (LOC283350), mRNA
 XM_208647 Homo sapiens hypothetical protein LOC283377 (LOC283377), mRNA
 XM_208658 Homo sapiens similar to Succinyl-CoA ligase [GDP-forming] beta-chain, mito
 XM_208667 Homo sapiens similar to HEE9341 (LOC283420), mRNA
 XM_208690 Homo sapiens similar to 40S ribosomal protein S28 (LOC283479), mRNA
 XM_208731 Homo sapiens chromosome 14 open reading frame 68 (C14orf68), mRNA
 XM_208746 Homo sapiens hypothetical protein LOC283578 (LOC283578), mRNA
 XM_208766 Homo sapiens KIAA0284 (KIAA0284), mRNA
 XM_208778 Homo sapiens hypothetical LOC283677 (LOC283677), mRNA
 XM_208798 Homo sapiens similar to hypothetical protein FLJ35785 (LOC283717), mRN/
 XM_208809 Homo sapiens similar to hypothetical protein D030069K18 (LOC283726), mF
 XM_208835 Homo sapiens similar to hypothetical protein FLJ36144 (LOC283767), mRN/
 XM_208847 Homo sapiens similar to testicular Metalloprotease-like, Disintegrin-like, Cyst
 XM_208859 Homo sapiens similar to ARHGAP21 protein (LOC283816), mRNA
 XM_208863 Homo sapiens similar to nuclear pore complex interacting protein (LOC34811
 XM_208887 Homo sapiens hypothetical protein LOC283871 (LOC283871), mRNA
 XM_208889 Homo sapiens similar to MGC9515 protein (LOC388240), mRNA
 XM_208891 Homo sapiens similar to nuclear pore complex interacting protein (LOC28381
 XM_208908 Homo sapiens hypothetical protein LOC283922 (LOC283922), mRNA
 XM_208927 Homo sapiens hypothetical protein FLJ36208 (FLJ36208), mRNA
 XM_208930 Homo sapiens similar to RIKEN cDNA 493051J11 (LOC283953), mRNA
 XM_208944 Homo sapiens hypothetical protein LOC283989 (LOC283989), mRNA

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XM_208990 Homo sapiens hypothetical LOC284067 (LOC284067), mRNA
 XM_208993 Homo sapiens Similar to hypothetical gene supported by AL050367; AK0229
 XM_209012 Homo sapiens similar to keratin 17 (LOC284089), mRNA
 XM_209041 Homo sapiens similar to KIAA1503 protein (LOC284158), mRNA
 XM_209073 Homo sapiens hypothetical protein LOC284207 (LOC284207), mRNA
 XM_209076 Homo sapiens similar to Ankyrin repeat domain protein 18A (LOC284232), r
 XM_209083 Homo sapiens similar to telomeric repeat binding factor 1 isoform 2; Telomer
 XM_209097 Homo sapiens similar to FLJ10101 protein (LOC284269), mRNA
 XM_209104 Homo sapiens similar to Placental thrombin inhibitor (Cytoplasmic antiprotein
 XM_209111 Homo sapiens hypothetical protein LOC284307 (LOC284307), mRNA
 XM_209138 Homo sapiens similar to LL5 beta (LOC388548), mRNA
 XM_209140 Homo sapiens hypothetical protein LOC284323 (LOC284323), mRNA
 XM_209145 Homo sapiens similar to RIKEN cDNA 0610012D14 (LOC284363), mRNA
 XM_209149 Homo sapiens similar to hypothetical protein FLJ31030 (LOC284318), mRNA/
 XM_209155 Homo sapiens hypothetical protein LOC284371 (LOC284371), mRNA
 XM_209163 Homo sapiens similar to solute carrier family 7, member 3; amino acid transp
 XM_209178 Homo sapiens similar to 60S ribosomal protein L10 (QM protein homolog) (L
 XM_209180 Homo sapiens similar to FKSG60 (LOC284397), mRNA
 XM_209187 Homo sapiens similar to BC022651 protein (LOC284417), mRNA
 XM_209196 Homo sapiens similar to HSPC323 (LOC284422), mRNA
 XM_209204 Homo sapiens hypothetical protein MGC26694 (MGC26694), mRNA
 XM_209227 Homo sapiens similar to KIAA0456 protein (LOC391087), mRNA
 XM_209234 Homo sapiens hypothetical protein DKFZp434E1410 (DKFZp434E1410), mF
 XM_209252 Homo sapiens similar to RIKEN cDNA 4930522H14 (LOC284548), mRNA
 XM_209337 Homo sapiens similar to coiled-coil-helix-coiled-coil-helix domain containing ;
 XM_209363 Homo sapiens protein tyrosine phosphatase, non-receptor type substrate 1-II
 XM_209384 Homo sapiens hypothetical LOC284859 (LOC284859), mRNA
 XM_209394 Homo sapiens hypothetical LOC284874 (LOC284874), mRNA
 XM_209408 Homo sapiens similar to bA436C9.2 (PUTATIVE novel protein similar to part
 XM_209423 Homo sapiens similar to bA395L14.5 (novel phosphoglucomutase like protei
 XM_209429 Homo sapiens similar to ARHQ protein (LOC284988), mRNA
 XM_209489 Homo sapiens similar to CG14853-PB (LOC285141), mRNA
 XM_209490 Homo sapiens hypothetical protein LOC285148 (LOC285148), mRNA
 XM_209500 Homo sapiens similar to 60S ribosomal protein L10 (QM protein homolog) (L
 XM_209505 Homo sapiens similar to KIAA0445 protein (LOC285188), mRNA
 XM_209509 Homo sapiens similar to RIKEN cDNA 0710001B24 (LOC285193), mRNA
 XM_209550 Homo sapiens similar to Carboxypeptidase N 83 kDa chain (Carboxypeptidase
 XM_209554 Homo sapiens similar to FSHD Region Gene 2 protein (LOC285299), mRNA
 XM_209559 Homo sapiens similar to MGC27169 protein (LOC285303), mRNA
 XM_209563 Homo sapiens hypothetical LOC285311 (LOC285311), mRNA
 XM_209569 Homo sapiens similar to ataxin-1 ubiquitin-like interacting protein (LOC2853
 XM_209579 Homo sapiens hypothetical protein LOC285346 (LOC285346), mRNA
 XM_209587 Homo sapiens similar to beta-1,4-mannosyltransferase; beta-1,4 mannosyltr
 XM_209604 Homo sapiens hypothetical LOC285423 (LOC285423), mRNA
 XM_209607 Homo sapiens hypothetical protein LOC285429 (LOC285429), mRNA
 XM_209612 Homo sapiens hypothetical protein LOC285440 (LOC285440), mRNA
 XM_209616 Homo sapiens similar to APACD protein (LOC285453), mRNA
 XM_209640 Homo sapiens hypothetical protein LOC285501 (LOC285501), mRNA
 XM_209643 Homo sapiens similar to hypothetical protein FLJ20035 (LOC285510), mRNA/
 XM_209655 Homo sapiens similar to beta-1,4-mannosyltransferase; beta-1,4 mannosyltr
 XM_209656 Homo sapiens hypothetical protein LOC285550 (LOC285550), mRNA
 XM_209668 Homo sapiens similar to RIKEN cDNA 1700007I06 (LOC285588), mRNA
 XM_209695 Homo sapiens similar to Chromosome-associated kinesin KIF4A (Chromok
 XM_209700 Homo sapiens similar to FLJ00052 protein (LOC285647), mRNA
 XM_209704 Homo sapiens similar to hypothetical protein (LOC285658), mRNA
 XM_209719 Homo sapiens hypothetical protein LOC285679 (LOC285679), mRNA
 XM_209728 Homo sapiens similar to C-terminal binding protein 2 isoform 2; ribeye (LOC

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XM_209741 Homo sapiens similar to Translationally controlled tumor protein (TCTP) (p23)
 XM_209763 Homo sapiens similar to FLJ00254 protein (LOC285770), mRNA
 XM_209824 Homo sapiens similar to matrilin 2 precursor (LOC285929), mRNA
 XM_209827 Homo sapiens similar to 40S ribosomal protein S26 (LOC285938), mRNA
 XM_209870 Homo sapiens similar to FLJ10408 protein (LOC286040), mRNA
 XM_209889 Homo sapiens similar to hypothetical protein (LOC286076), mRNA
 XM_209893 Homo sapiens similar to TBP-associated factor 15 isoform 1; TAF15 RNA po
 XM_209902 Homo sapiens zinc finger protein 252 (ZNF252), mRNA
 XM_209910 Homo sapiens similar to chromosome 15 open reading frame 2 (LOC286129
 XM_209913 Homo sapiens similar to ring finger protein 5 (LOC286140), mRNA
 XM_209918 Homo sapiens similar to RIKEN cDNA 4930533G20 (LOC286151), mRNA
 XM_209920 Homo sapiens similar to mCBP (LOC286157), mRNA
 XM_209936 Homo sapiens similar to RIKEN cDNA 1700011J18 (LOC286187), mRNA
 XM_209941 Homo sapiens hypothetical protein LOC286207 (LOC286207), mRNA
 XM_209955 Homo sapiens similar to beta-tubulin 4Q (LOC286222), mRNA
 XM_209967 Homo sapiens similar to NAD-dependent malic enzyme, mitochondrial precu
 XM_210022 Homo sapiens GTPase activating RANGAP domain-like 1 (GARNL1), mRNA
 XM_210035 Homo sapiens similar to putative UST1-like organic anion transporter (LOC2
 XM_210042 Homo sapiens similar to mitochondrial ribosome recycling factor isoform 1 (L
 XM_210048 Homo sapiens hypothetical protein LOC286436 (LOC286436), mRNA
 XM_210054 Homo sapiens similar to dJ341D10.1 (novel protein) (LOC286453), mRNA
 XM_210062 Homo sapiens ras-related C3 botulinum toxin substrate family member 4 (R
 XM_210082 Homo sapiens similar to 40S ribosomal protein S26 (LOC286513), mRNA
 XM_210094 Homo sapiens similar to 40S ribosomal protein S26 (LOC286539), mRNA
 XM_210119 Homo sapiens similar to Olfactory receptor 5AT1 (LOC284532), mRNA
 XM_210168 Homo sapiens similar to Olfactory receptor 4C13 (LOC283092), mRNA
 XM_210169 Homo sapiens similar to Olfactory receptor 4C12 (LOC283093), mRNA
 XM_210180 Homo sapiens similar to odorant receptor HOR3beta2 (LOC283110), mRNA
 XM_210181 Homo sapiens similar to odorant receptor HOR3beta1 (LOC283111), mRNA
 XM_210184 Homo sapiens similar to tripartite motif-containing 43 (LOC283117), mRNA
 XM_210186 Homo sapiens similar to Olfactory receptor 8B4 (LOC283162), mRNA
 XM_210193 Homo sapiens similar to Olfactory receptor 9G4 (LOC283189), mRNA
 XM_210196 Homo sapiens similar to seven transmembrane helix receptor (LOC283193)
 XM_210242 Homo sapiens similar to olfactory receptor GA_x6K02T2PULF-11031172-111
 XM_210282 Homo sapiens similar to Olfactory receptor 4K5 (LOC283621), mRNA
 XM_210324 Homo sapiens similar to Vacuolar ATP synthase 16 kDa proteolipid subunit (n
 XM_210334 Homo sapiens similar to 60S ribosomal protein L29 (Cell surface heparin bin
 XM_210365 Homo sapiens similar to Ribosomal protein L24-like (LOC284288), mRNA
 XM_210382 Homo sapiens similar to Olfactory receptor 221 (LOC284383), mRNA
 XM_210394 Homo sapiens similar to Contains similarity to extensin (atExt1) from Arabido
 XM_210400 Homo sapiens similar to autoantigen NOR-90 (LOC285031), mRNA
 XM_210411 Homo sapiens similar to Oligophrenin 1 (LOC285101), mRNA
 XM_210501 Homo sapiens similar to HSPC182 protein (LOC286528), mRNA
 XM_210515 Homo sapiens similar to testis specific protein, Y-linked (LOC286561), mRNA
 XM_210543 Homo sapiens similar to SPBPJ4664.02 (LOC285253), mRNA
 XM_210557 Homo sapiens similar to heterochromatin-specific nonhistone protein (LOC2
 XM_210559 Homo sapiens similar to Retinoic acid receptor beta (RAR-beta) (RAR-epsilo
 XM_210562 Homo sapiens hypothetical protein LOC285335 (LOC285335), mRNA
 XM_210576 Homo sapiens similar to RIKEN cDNA 4921517D21 (LOC285405), mRNA
 XM_210581 Homo sapiens claudin 22 (CLDN22), mRNA
 XM_210613 Homo sapiens similar to Transcription initiation factor TFIID 28 kDa subunit (n
 XM_210642 Homo sapiens similar to Transcription initiation factor TFIID 28 kDa subunit (n
 XM_210752 Homo sapiens similar to Olfactory receptor 13C9 (LOC286362), mRNA
 XM_210755 Homo sapiens similar to Olfactory receptor 13D1 (LOC286365), mRNA
 XM_210787 Homo sapiens similar to Olfactory receptor 10H5 (LOC284433), mRNA
 XM_210826 Homo sapiens similar to Wiskott-Aldrich syndrome protein family member 4 (n
 XM_210856 Homo sapiens similar to Ras suppressor protein 1 (LOC283029), mRNA

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XM_210860 Homo sapiens hypothetical LOC283034 (LOC283034), mRNA
 XM_210876 Homo sapiens hypothetical LOC283065 (LOC283065), mRNA
 XM_210906 Homo sapiens hypothetical LOC283166 (LOC283166), mRNA
 XM_210908 Homo sapiens similar to RIKEN cDNA A930008A22 (LOC283157), mRNA
 XM_211009 Homo sapiens hypothetical LOC283389 (LOC283389), mRNA
 XM_211028 Homo sapiens hypothetical protein LOC283403 (LOC283403), mRNA
 XM_211040 Homo sapiens hypothetical LOC283440 (LOC283440), mRNA
 XM_211079 Homo sapiens hypothetical LOC283530 (LOC283530), mRNA
 XM_211086 Homo sapiens hypothetical LOC283553 (LOC283553), mRNA
 XM_211088 Homo sapiens hypothetical LOC283604 (LOC283604), mRNA
 XM_211089 Homo sapiens hypothetical LOC283586 (LOC283586), mRNA
 XM_211090 Homo sapiens hypothetical protein LOC283587 (LOC283587), mRNA
 XM_211092 Homo sapiens hypothetical LOC283583 (LOC283583), mRNA
 XM_211108 Homo sapiens hypothetical LOC283584 (LOC283584), mRNA
 XM_211174 Homo sapiens hypothetical LOC283710 (LOC283710), mRNA
 XM_211197 Homo sapiens hypothetical LOC283780 (LOC283780), mRNA
 XM_211244 Homo sapiens hypothetical LOC283895 (LOC283895), mRNA
 XM_211246 Homo sapiens similar to BTG3 associated nuclear protein isoform a; BANP 1
 XM_211251 Homo sapiens hypothetical LOC283902 (LOC283902), mRNA
 XM_211287 Homo sapiens hypothetical protein LOC283999 (LOC283999), mRNA
 XM_211291 Homo sapiens hypothetical protein LOC283994 (LOC283994), mRNA
 XM_211305 Homo sapiens hypothetical protein LOC284021 (LOC284021), mRNA
 XM_211339 Homo sapiens hypothetical LOC284120 (LOC284120), mRNA
 XM_211345 Homo sapiens hypothetical LOC284134 (LOC284134), mRNA
 XM_211367 Homo sapiens hypothetical LOC284184 (LOC284184), mRNA
 XM_211403 Homo sapiens CD33 antigen-like 3 (CD33L3), mRNA
 XM_211408 Homo sapiens hypothetical LOC284260 (LOC284260), mRNA
 XM_211413 Homo sapiens hypothetical LOC284275 (LOC284275), mRNA
 XM_211422 Homo sapiens hypothetical LOC284303 (LOC284303), mRNA
 XM_211432 Homo sapiens hypothetical LOC284321 (LOC284321), mRNA
 XM_211447 Homo sapiens hypothetical LOC284409 (LOC284409), mRNA
 XM_211460 Homo sapiens hypothetical protein LOC284434 (LOC284434), mRNA
 XM_211509 Homo sapiens hypothetical LOC284527 (LOC284527), mRNA
 XM_211518 Homo sapiens hypothetical LOC284555 (LOC284555), mRNA
 XM_211529 Homo sapiens hypothetical protein LOC284591 (LOC284591), mRNA
 XM_211557 Homo sapiens hypothetical LOC284623 (LOC284623), mRNA
 XM_211573 Homo sapiens hypothetical LOC284646 (LOC284646), mRNA
 XM_211627 Homo sapiens hypothetical LOC284754 (LOC284754), mRNA
 XM_211694 Homo sapiens hypothetical LOC284931 (LOC284931), mRNA
 XM_211707 Homo sapiens hypothetical LOC284968 (LOC284968), mRNA
 XM_211736 Homo sapiens hypothetical protein LOC285016 (LOC285016), mRNA
 XM_211749 Homo sapiens hypothetical LOC285047 (LOC285047), mRNA
 XM_211764 Homo sapiens hypothetical LOC285095 (LOC285095), mRNA
 XM_211768 Homo sapiens hypothetical LOC285110 (LOC285110), mRNA
 XM_211805 Homo sapiens hypothetical protein LOC285205 (LOC285205), mRNA
 XM_211816 Homo sapiens hypothetical LOC285248 (LOC285248), mRNA
 XM_211837 Homo sapiens hypothetical LOC285307 (LOC285307), mRNA
 XM_211843 Homo sapiens hypothetical protein LOC285326 (LOC285326), mRNA
 XM_211853 Homo sapiens similar to hypothetical protein MG11009.4 (LOC285344), MR1
 XM_211858 Homo sapiens hypothetical protein FLJ12205 (LOC285359), mRNA
 XM_211871 Homo sapiens hypothetical LOC285382 (LOC285382), mRNA
 XM_211896 Homo sapiens hypothetical LOC285435 (LOC285435), mRNA
 XM_211908 Homo sapiens similar to unc-93 homolog B1; unc93 (C.elegans) homolog B;
 XM_211923 Homo sapiens hypothetical LOC285509 (LOC285509), mRNA
 XM_211983 Homo sapiens hypothetical LOC285694 (LOC285694), mRNA
 XM_211988 Homo sapiens hypothetical LOC285711 (LOC285711), mRNA
 XM_211995 Homo sapiens hypothetical LOC285721 (LOC285721), mRNA

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XM_212013 Homo sapiens hypothetical LOC285777 (LOC285777), mRNA
 XM_212022 Homo sapiens hypothetical LOC285793 (LOC285793), mRNA
 XM_212061 Homo sapiens similar to THAP domain containing 5 (LOC285872), mRNA
 XM_212067 Homo sapiens hypothetical LOC285890 (LOC285890), mRNA
 XM_212094 Homo sapiens hypothetical LOC285919 (LOC285919), mRNA
 XM_212123 Homo sapiens hypothetical LOC285995 (LOC285995), mRNA
 XM_212162 Homo sapiens similar to CG3104-PA (LOC286080), mRNA
 XM_212170 Homo sapiens hypothetical protein LOC286094 (LOC286094), mRNA
 XM_212238 Homo sapiens hypothetical protein LOC286235 (LOC286235), mRNA
 XM_212241 Homo sapiens DKFZP434M131 protein (DKFZP434M131), mRNA
 XM_212319 Homo sapiens hypothetical LOC286441 (LOC286441), mRNA
 XM_212326 Homo sapiens hypothetical LOC286478 (LOC286478), mRNA
 XM_212581 Homo sapiens zinc finger protein 311 (ZNF311), mRNA
 XM_290185 Homo sapiens similar to hypothetical protein MGC5560 (LOC338598), mRNA/
 XM_290225 Homo sapiens similar to pote protein; Expressed in prostate, ovary, testis, ar
 XM_290331 Homo sapiens gamma-glutamyltransferase 2 (GGT2), mRNA
 XM_290342 Homo sapiens similar to immunoglobulin superfamily, member 3; immunoglo
 XM_290345 Homo sapiens similar to eukaryotic translation initiation factor 3, subunit 5 e
 XM_290351 Homo sapiens similar to Nedd4-like ubiquitin-protein ligase WWP1 (WW do
 XM_290385 Homo sapiens similar to solute carrier family 29 (nucleoside transporters), m
 XM_290389 Homo sapiens similar to solute carrier family 29 (nucleoside transporters), m
 XM_290401 Homo sapiens hypothetical protein LOC340318 (LOC340318), mRNA
 XM_290462 Homo sapiens KIAA1674 (KIAA1674), mRNA
 XM_290463 Homo sapiens family with sequence similarity 22, member A (FAM22A), mRf
 XM_290482 Homo sapiens hypothetical protein FLJ10824 (FLJ10824), mRNA
 XM_290501 Homo sapiens hypothetical LOC338661 (LOC338661), mRNA
 XM_290502 Homo sapiens KIAA1030 protein (KIAA1030), mRNA
 XM_290506 Homo sapiens splicing factor 3b, subunit 2, 145kDa (SF3B2), mRNA
 XM_290509 Homo sapiens hypothetical protein LOC338692 (LOC338692), mRNA
 XM_290516 Homo sapiens myotonic dystrophy protein kinase like protein (HSMDPKIN), r
 XM_290517 Homo sapiens KIAA0404 protein (KIAA0404), mRNA
 XM_290527 Homo sapiens ubiquitin specific protease 35 (USP35), mRNA
 XM_290536 Homo sapiens CTD-binding SR-like protein rA9 (KIAA1542), mRNA
 XM_290546 Homo sapiens KIAA0830 protein (KIAA0830), mRNA
 XM_290547 Homo sapiens similar to Oligophrenin 1 (LOC338734), mRNA
 XM_290552 Homo sapiens cyclic nucleotide gated channel alpha 4 (CNGA4), mRNA
 XM_290558 Homo sapiens similar to C1q-related factor precursor (LOC338751), mRNA
 XM_290599 Homo sapiens glutamate receptor interacting protein 1 (GRIP1), mRNA
 XM_290579 Homo sapiens hypothetical protein LOC338797 (LOC338797), mRNA
 XM_290592 Homo sapiens huntingtin interacting protein-1-related (HIP1R), mRNA
 XM_290597 Homo sapiens hypothetical protein LOC283464 (LOC283464), mRNA
 XM_290598 Homo sapiens dendrin (DDN), mRNA
 XM_290600 Homo sapiens similar to Reticulocalbin 1 precursor (LOC338851), mRNA
 XM_290609 Homo sapiens hypothetical LOC338914 (LOC338914), mRNA
 XM_290615 Homo sapiens hypothetical protein DKFZp762F0713 (DKFZp762F0713), mR
 XM_290626 Homo sapiens similar to chromosome 1 open reading frame 36 (LOC338934
 XM_290629 Homo sapiens chromosome 14 open reading frame 78 (C14orf78), mRNA
 XM_290631 Homo sapiens glucuronyl C5-epimerase (GLCE), mRNA
 XM_290645 Homo sapiens hypothetical protein DKFZp434J0617 (DKFZp434J0617), mR
 XM_290660 Homo sapiens similar to HP95 (LOC339003), mRNA
 XM_290667 Homo sapiens KIAA0350 protein (KIAA0350), mRNA
 XM_290670 Homo sapiens KIAA0220 protein (KIAA0220), mRNA
 XM_290671 Homo sapiens hypothetical protein LOC339047 (LOC339047), mRNA
 XM_290684 Homo sapiens DKFZP434I216 protein (DKFZP434I216), mRNA
 XM_290702 Homo sapiens similar to Myo16 protein (LOC339088), mRNA
 XM_290704 Homo sapiens hypothetical protein FLJ12270 (FLJ12270), mRNA
 XM_290712 Homo sapiens hypothetical protein MGC46336 (MGC46336), mRNA

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XM_290714 Homo sapiens RAB43, member RAS oncogene family (RAB43), mRNA
 XM_290722 Homo sapiens similar to RIKEN cDNA 2810003J06 (LOC339123), mRNA
 XM_290732 Homo sapiens KIAA1917 protein (KIAA1917), mRNA
 XM_290734 Homo sapiens similar to ataxin 2 binding protein 1 isoform gamma; hexanobi
 XM_290737 Homo sapiens KIAA1871 protein (KIAA1871), mRNA
 XM_290743 Homo sapiens similar to hypothetical protein FLJ36492 (LOC339184), mRNA/
 XM_290755 Homo sapiens hypothetical protein FLJ35848 (FLJ35848), mRNA
 XM_290758 Homo sapiens KIAA0553 protein (KIAA0553), mRNA
 XM_290768 Homo sapiens KIAA1554 protein (KIAA1554), mRNA
 XM_290777 Homo sapiens hypothetical protein LOC339231 (LOC339231), mRNA
 XM_290780 Homo sapiens similar to Envoiplakin (210 kDa paraneoplastic pemphigus ant
 XM_290781 Homo sapiens similar to Keratin, type I cytoskeletal 14 (Cytokeratin 14) (K14
 XM_290782 Homo sapiens similar to Keratin, type I cytoskeletal 16 (Cytokeratin 16) (K16
 XM_290786 Homo sapiens similar to endozepine-like protein (LOC339253), mRNA
 XM_290793 Homo sapiens kinase suppressor of ras (KSR), mRNA
 XM_290795 Homo sapiens hypothetical protein DKFZp667M2411 (DKFZp667M2411), ml
 XM_290799 Homo sapiens KIAA1501 protein (KIAA1501), mRNA
 XM_290809 Homo sapiens TAF4b RNA polymerase II, TATA box binding protein (TBP)-a
 XM_290811 Homo sapiens KIAA1713 protein (KIAA1713), mRNA
 XM_290817 Homo sapiens hypothetical protein FLJ34907 (FLJ34907), mRNA
 XM_290818 Homo sapiens Spir-1 protein (Spir-1), mRNA
 XM_290820 Homo sapiens hypothetical protein FLJ10211 (FLJ10211), mRNA
 XM_290822 Homo sapiens hypothetical protein LOC284367 (LOC284367), mRNA
 XM_290829 Homo sapiens homolog of Drosophila Intersex (Intersex), mRNA
 XM_290831 Homo sapiens hypothetical protein LOC339321 (LOC339321), mRNA
 XM_290835 Homo sapiens zinc finger protein 181 (FHZ181) (ZNF181), mRNA
 XM_290838 Homo sapiens hypothetical protein LOC339324 (LOC339324), mRNA
 XM_290842 Homo sapiens leucine rich repeat and fibronectin type III domain containing
 XM_290848 Homo sapiens hypothetical protein LOC339344 (LOC339344), mRNA
 XM_290850 Homo sapiens glutamate receptor, ionotropic, N-methyl-D-aspartate 3B (GRI
 XM_290854 Homo sapiens G protein-coupled receptor 108 (GPR108), mRNA
 XM_290865 Homo sapiens similar to Zinc finger protein 93 (Zinc finger protein HTF34) (L
 XM_290866 Homo sapiens similar to 4930572L20Rik protein (LOC339377), mRNA
 XM_290867 Homo sapiens RalGEF-like protein 3, mouse homolog (RGL3), mRNA
 XM_290872 Homo sapiens similar to Pyruvate kinase, M2 isozyme (LOC339395), mRNA
 XM_290902 Homo sapiens hypothetical protein LOC339448 (LOC339448), mRNA
 XM_290922 Homo sapiens hypothetical protein MGC27277 (MGC27277), mRNA
 XM_290923 Homo sapiens KIAA1639 protein (KIAA1639), mRNA
 XM_290925 Homo sapiens hypothetical LOC339494 (LOC339494), mRNA
 XM_290927 Homo sapiens similar to bA47615.3 (novel protein similar to septin) (LOC33:
 XM_290936 Homo sapiens hypothetical protein MGC35030 (MGC35030), mRNA
 XM_290941 Homo sapiens prion protein interacting protein (PRNPiP), mRNA
 XM_290944 Homo sapiens KIAA0842 protein (KIAA0842), mRNA
 XM_290948 Homo sapiens hypothetical protein LOC343071 (LOC343071), mRNA
 XM_290949 Homo sapiens similar to hypothetical protein (LOC339553), mRNA
 XM_290972 Homo sapiens novel C3HC4 type Zinc finger (ring finger) (FLJ12747), mRNA
 XM_290973 Homo sapiens solute carrier family 35, member E4 (SLC35E4), mRNA
 XM_290985 Homo sapiens hypothetical protein LOC339692 (LOC339692), mRNA
 XM_290994 Homo sapiens similar to speckle-type POZ protein (LOC339744), mRNA
 XM_291001 Homo sapiens myosin VIIb (MYO7B), mRNA
 XM_291005 Homo sapiens similar to 25-hydroxyvitamin D-1 alpha hydroxylase, mitochon
 XM_291007 Homo sapiens hypothetical protein LOC339766 (LOC339766), mRNA
 XM_291015 Homo sapiens likely homolog of rat kinase D-interacting substance of 220 kE
 XM_291016 Homo sapiens similar to RIKEN cDNA 1700093K21 (LOC339804), mRNA
 XM_291017 Homo sapiens similar to PRO1094 (LOC339793), mRNA
 XM_291019 Homo sapiens hypothetical protein FLJ13305 (FLJ13305), mRNA
 XM_291020 Homo sapiens KIAA2012 protein (LOC339809), mRNA

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XM_291028 Homo sapiens hypothetical protein DKFZp434A128 (DKFZp434A128), mRNA
 XM_291054 Homo sapiens similar to beta-1,4-mannosyltransferase; beta-1,4-mannosyltri
 XM_291055 Homo sapiens KIAA1268 protein (KIAA1268), mRNA
 XM_291057 Homo sapiens cytoplasmic linker associated protein 2 (CLASP2), mRNA
 XM_291059 Homo sapiens hypothetical protein LOC339896 (LOC339896), mRNA
 XM_291062 Homo sapiens similar to K06A9.1b.p (KIAA2018), mRNA
 XM_291063 Homo sapiens hypothetical protein LOC339903 (LOC339903), mRNA
 XM_291064 Homo sapiens KIAA0540 protein (KIAA0540), mRNA
 XM_291074 Homo sapiens hypothetical protein FLJ33674 (FLJ33674), mRNA
 XM_291075 Homo sapiens hypothetical LOC339926 (LOC339926), mRNA
 XM_291077 Homo sapiens hypothetical protein BC009862 (LOC90113), mRNA
 XM_291080 Homo sapiens KIAA0804 protein (KIAA0804), mRNA
 XM_291085 Homo sapiens KIAA1204 protein (CDGAP), mRNA
 XM_291088 Homo sapiens similar to hypothetical protein SB153 isoform 1 (LOC339944),
 XM_291095 Homo sapiens similar to glutamate receptor, ionotropic, N-methyl D-aspartate
 XM_291099 Homo sapiens similar to Hypothetical protein MGC38937 (LOC339977), mR
 XM_291105 Homo sapiens transcriptional adaptor 2 (ADA2 homolog, yeast)-beta (MGC2
 XM_291106 Homo sapiens KIAA0232 gene product (KIAA0232), mRNA
 XM_291111 Homo sapiens G protein-coupled receptor 125 (GPR125), mRNA
 XM_291117 Homo sapiens similar to succinate dehydrogenase flavoprotein subunit (LOC
 XM_291120 Homo sapiens hypothetical protein LOC340024 (LOC340024), mRNA
 XM_291128 Homo sapiens KIAA1311 protein (KIAA1311), mRNA
 XM_291137 Homo sapiens similar to hypothetical protein D630003M2.1 (LOC389256), m
 XM_291139 Homo sapiens similar to RIKEN cDNA 9330196J05 (LOC340075), mRNA
 XM_291141 Homo sapiens KIAA0303 protein (KIAA0303), mRNA
 XM_291142 Homo sapiens hypothetical protein BC014311 (LOC115548), mRNA
 XM_291144 Homo sapiens similar to bA110H4.2 (similar to membrane protein) (LOC340
 XM_291154 Homo sapiens similar to hypothetical protein D730019B10 (LOC340152), m
 XM_291159 Homo sapiens ion transporter protein (DKFZP434F011), mRNA
 XM_291181 Homo sapiens similar to developmental pluripotency associated 5; embryone
 XM_291189 Homo sapiens similar to precursor peptide (LOC340204), mRNA
 XM_291170 Homo sapiens similar to Hypothetical protein KIAA0036 (LOC340192), mRN
 XM_291181 Homo sapiens similar to LINE-1 REVERSE TRANSCRIPTASE HOMOLOG (I
 XM_291200 Homo sapiens similar to CAGL79 (LOC340221), mRNA
 XM_291202 Homo sapiens zinc finger protein 479 (ZNF479), mRNA
 XM_291204 Homo sapiens hypothetical LOC340228 (LOC340228), mRNA
 XM_291208 Homo sapiens similar to hypothetical ZNF-like protein (LOC340246), mRNA
 XM_291222 Homo sapiens DKFZP586J0619 protein (DKFZP586J0619), mRNA
 XM_291223 Homo sapiens myosin IG (MYO1G), mRNA
 XM_291241 Homo sapiens intracellular membrane-associated calcium-independent phos
 XM_291247 Homo sapiens similar to Piccolo protein (Aczonin) (LOC389530), mRNA
 XM_291253 Homo sapiens KIAA0146 protein (KIAA0146), mRNA
 XM_291261 Homo sapiens zinc finger protein 517 (ZNF517), mRNA
 XM_291262 Homo sapiens zinc finger protein 251 (ZNF251), mRNA
 XM_291266 Homo sapiens 5-oxoprolinase (ATP-hydrolysing) (OPLAH), mRNA
 XM_291268 Homo sapiens glutamate receptor, ionotropic, N-methyl D-aspartate-associat
 XM_291269 Homo sapiens hypothetical protein KIAA1875 (KIAA1875), mRNA
 XM_291270 Homo sapiens similar to unmaed protein product (LOC340393), mRNA
 XM_291277 Homo sapiens hypothetical protein DKFZp761P0423 (DKFZp761P0423), mR
 XM_291282 Homo sapiens hypothetical protein LOC157697 (LOC157697), mRNA
 XM_291291 Homo sapiens KIAA0725 protein (KIAA0725), mRNA
 XM_291314 Homo sapiens F-box only protein 10 (FBXO10), mRNA
 XM_291315 Homo sapiens KIAA1815 (KIAA1815), mRNA
 XM_291321 Homo sapiens similar to Nance-Horan syndrome protein (LOC340527), mRN
 XM_291322 Homo sapiens KIAA2001 protein (KIAA2001), mRNA
 XM_291326 Homo sapiens KIAA2022 protein (KIAA2022), mRNA
 XM_291334 Homo sapiens similar to hypothetical protein MGC15737 (LOC340543), mRN

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XM_291335 Homo sapiens hypothetical protein LOC340542 (LOC340542), mRNA
 XM_291339 Homo sapiens hypothetical protein LOC340562 (LOC340562), mRNA
 XM_291344 Homo sapiens hypothetical protein FLJ12649 (FLJ12649), mRNA
 XM_291345 Homo sapiens KIAA0522 protein (KIAA0522), mRNA
 XM_291346 Homo sapiens similar to carbonic anhydrase VB, mitochondrial precursor; ca
 XM_291378 Homo sapiens similar to C4b-binding protein alpha chain precursor (C4bp) (f
 XM_291387 Homo sapiens similar to DAAT9248 (LOC339398), mRNA
 XM_291392 Homo sapiens similar to Arylacetamide deacetylase (AADAC) (LOC343066),
 XM_291394 Homo sapiens hypothetical protein LOC343068 (LOC343068), mRNA
 XM_291395 Homo sapiens similar to Heterogeneous nuclear ribonucleoprotein C-like dJf
 XM_291396 Homo sapiens similar to Hypothetical protein DJ845O24.2 (LOC400735), mF
 XM_291400 Homo sapiens similar to chromosome 14 open reading frame 24 (LOC34308
 XM_291419 Homo sapiens G protein-coupled receptor 153 (GPR153), mRNA
 XM_291428 Homo sapiens similar to ribosomal protein L26 (LOC343153), mRNA
 XM_291435 Homo sapiens similar to Olfactory receptor 6F1 (LOC343169), mRNA
 XM_291436 Homo sapiens similar to Olfactory receptor 5AY1 (LOC343170), mRNA
 XM_291437 Homo sapiens olfactory receptor, family 1, subfamily C, member 1 (OR1C1),
 XM_291438 Homo sapiens similar to seven transmembrane helix receptor (LOC343171),
 XM_291439 Homo sapiens similar to seven transmembrane helix receptor (LOC343172),
 XM_291441 Homo sapiens similar to Olfactory receptor 2T3 (LOC343173), mRNA
 XM_291464 Homo sapiens similar to KIAA0433 protein (LOC343221), mRNA
 XM_291485 Homo sapiens similar to Myosin-binding protein H (MyBP-H) (H-protein) (LOC
 XM_291508 Homo sapiens similar to Farnesyl pyrophosphate synthetase (FPP synthetase
 XM_291533 Homo sapiens similar to basic transcription factor 3 (LOC343363), mRNA
 XM_291543 Homo sapiens similar to KIAA0454 protein (LOC343381), mRNA
 XM_291544 Homo sapiens similar to peptidyl-Pro cis trans isomerase (LOC343384), mRf
 XM_291548 Homo sapiens similar to Olfactory receptor 10R2 (LOC343406), mRNA
 XM_291569 Homo sapiens similar to IFGP8 (LOC343413), mRNA
 XM_291577 Homo sapiens similar to RIKEN cDNA 1810009O10 (LOC339521), mRNA
 XM_291584 Homo sapiens similar to Phosphorylase B kinase alpha regulatory chain, ske
 XM_291607 Homo sapiens similar to dJ1182A14.5.1 (novel gene (isoform 1)) (LOC3435C
 XM_291623 Homo sapiens similar to hypothetical protein 4833401D15 (LOC343521), mF
 XM_291625 Homo sapiens similar to Hypothetical protein DJ845O24.2 (LOC374947), mF
 XM_291627 Homo sapiens similar to oral cancer overexpressed 2; transmembrane protel
 XM_291638 Homo sapiens hypothetical protein LOC343070 (LOC343070), mRNA
 XM_291643 Homo sapiens similar to Ig kappa chain (LOC339562), mRNA
 XM_291645 Homo sapiens similar to Olfactory receptor 2T5 (LOC343563), mRNA
 XM_291663 Homo sapiens similar to bA304I5.1 (novel lipase) (LOC340654), mRNA
 XM_291671 Homo sapiens hypothetical protein LOC282996 (LOC282996), mRNA
 XM_291697 Homo sapiens similar to RIKEN cDNA A930010E21 (LOC340745), mRNA
 XM_291698 Homo sapiens similar to Rpl7a protein (LOC340749), mRNA
 XM_291704 Homo sapiens similar to double homeobox, 4; double homeobox protein 4 (L
 XM_291716 Homo sapiens similar to NK-type homeobox (LOC340784), mRNA
 XM_291723 Homo sapiens protein RAKc (LOC340811), mRNA
 XM_291725 Homo sapiens alpha 2,8-sialyltransferase (ST8SIA-VI), mRNA
 XM_291726 Homo sapiens similar to protein of unknown function (LOC340843), mRNA
 XM_291729 Homo sapiens TAF3 RNA polymerase II, TATA box binding protein (TBP)-as
 XM_291741 Homo sapiens dual specificity phosphatase and pro isomerase domain conte
 XM_291745 Homo sapiens similar to 40S ribosomal protein S26 (LOC338611), mRNA
 XM_291757 Homo sapiens similar to protein of unknown function (LOC340893), mRNA
 XM_291763 Homo sapiens similar to bA508N22.1 (HSPC025) (LOC340947), mRNA
 XM_291767 Homo sapiens similar to hypothetical protein FLJ10213 (LOC340900), mRN/
 XM_291770 Homo sapiens similar to breast cancer antigen NY-BR-1 (LOC340913), mRN
 XM_291771 Homo sapiens similar to HSPC132 (LOC338617), mRNA
 XM_291786 Homo sapiens similar to Phosphorylase B kinase gamma catalytic chain, ske
 XM_291793 Homo sapiens similar to Chain A, Crystal Structure Of The Radixin Fern Dor
 XM_291806 Homo sapiens similar to seven transmembrane helix receptor (LOC340980),

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XM_291816 Homo sapiens otogelin (OTOG), mRNA
 XM_291838 Homo sapiens similar to Olfactory receptor 8D4 (LOC338662), mRNA
 XM_291857 Homo sapiens similar to ENSANGP00000020885 (LOC341098), mRNA
 XM_291859 Homo sapiens olfactory receptor, family 5, subfamily F, member 1 (OR5F1),
 XM_291862 Homo sapiens similar to Olfactory receptor 5AP2 (LOC338675), mRNA
 XM_291885 Homo sapiens similar to Tryptophanyl-tRNA synthetase (Tryptophan-tRNA II
 XM_291892 Homo sapiens similar to olfactory receptor MOR101-1 (LOC341152), mRNA
 XM_291924 Homo sapiens similar to Olfactory receptor 4F3 (LOC338718), mRNA
 XM_291943 Homo sapiens similar to Elongation factor 1-alpha 2 (EF-1-alpha-2) (Elongati
 XM_291947 Homo sapiens hephaestin-like (LOC341208), mRNA
 XM_291974 Homo sapiens similar to hypothetical protein FLJ33167 (LOC338750), mRNA/
 XM_291977 Homo sapiens similar to Olfactory receptor 52L1 (LOC338751), mRNA
 XM_291980 Homo sapiens similar to Olfactory receptor 2AG1 (HT3) (LOC338755), mRNA/
 XM_291981 Homo sapiens similar to hP4 olfactory receptor (LOC341276), mRNA
 XM_291986 Homo sapiens olfactory receptor, family 10, subfamily A, member 3 (OR10A:
 XM_291989 Homo sapiens similar to hypothetical protein (LOC338756), mRNA
 XM_291991 Homo sapiens similar to protein tyrosine phosphatase, receptor type, Q isofo
 XM_292012 Homo sapiens similar to Polyadenylate-binding protein 1 (Poly(A)-binding pr
 XM_292021 Homo sapiens similar to hypothetical protein (LOC341346), mRNA
 XM_292023 Homo sapiens similar to ribosomal protein L31 (LOC341356), mRNA
 XM_292027 Homo sapiens similar to Dag1-prov protein (LOC341370), mRNA
 XM_292029 Homo sapiens hypothetical LOC341371 (LOC341371), mRNA
 XM_292035 Homo sapiens similar to olfactory specific medium-chain acyl CoA synthetase
 XM_292046 Homo sapiens similar to ribosomal protein L31 (LOC341412), mRNA
 XM_292048 Homo sapiens similar to Heat shock factor protein 2 (HSF 2) (Heat shock tra
 XM_292049 Homo sapiens similar to olfactory receptor MOR114-1 (LOC341416), mRNA
 XM_292051 Homo sapiens similar to Olfactory receptor 6C4 (LOC341418), mRNA
 XM_292064 Homo sapiens similar to fertilin alpha subunit (LOC338792), mRNA
 XM_292085 Homo sapiens similar to peptidyl-Pro cis trans isomerase (LOC341457), mR
 XM_292093 Homo sapiens liver-specific organic anion transporter 3 (LST-3), mRNA
 XM_292098 Homo sapiens similar to hypothetical protein (LOC341461), mRNA
 XM_292109 Homo sapiens similar to 60S ribosomal protein L23a (LOC341511), mRNA
 XM_292122 Homo sapiens similar to neurofilament-like protein (LOC338829), mRNA
 XM_292136 Homo sapiens similar to seven transmembrane helix receptor (LOC341568),
 XM_292150 Homo sapiens similar to Serine/threonine-protein kinase Nek1 (NimA-related
 XM_292184 Homo sapiens similar to immune-responsive gene 1 (LOC341720), mRNA
 XM_292193 Homo sapiens hypothetical protein DKFZp686J0811 (DKFZp686J0811), mR
 XM_292197 Homo sapiens similar to bA215B13.2 (fumarate hydratase (FH) pseudogene,
 XM_292210 Homo sapiens similar to ribosomal protein S12 (LOC338870), mRNA
 XM_292225 Homo sapiens similar to putative pancreatic ribonuclease (LOC338879), mR
 XM_292227 Homo sapiens similar to Olfactory receptor 6S1 (LOC341799), mRNA
 XM_292280 Homo sapiens solute carrier family 35, member F4 (SLC35F4), mRNA
 XM_292301 Homo sapiens similar to developmental pluripotency associated 5; embryone
 XM_292357 Homo sapiens similar to Golgi autoantigen, golgin subfamily A member 6 (Gr
 XM_292384 Homo sapiens similar to zinc finger protein 29 (LOC342132), mRNA
 XM_292389 Homo sapiens similar to ATP-binding cassette, sub-family B, member 10, mi
 XM_292394 Homo sapiens similar to hypothetical protein FLJ35785 (LOC342167), mRNA/
 XM_292468 Homo sapiens similar to hypothetical protein (LOC342293), mRNA
 XM_292503 Homo sapiens similar to hypothetical protein (LOC342355), mRNA
 XM_292504 Homo sapiens similar to hypothetical protein FLJ35867 (LOC342357), mRNA/
 XM_292512 Homo sapiens similar to Ataxin-1 (Spinocerebellar ataxia type 1 protein hom
 XM_292527 Homo sapiens similar to Hypothetical protein MGC67567 (LOC342409), mR
 XM_292562 Homo sapiens similar to c439A6.1 (novel protein similar to heparan sulfate (c
 XM_292573 Homo sapiens similar to testicular Metalloprotease-like, Disintegrin-like, Cyst
 XM_292596 Homo sapiens similar to peptidyl-Pro cis trans isomerase (LOC342541), mR
 XM_292624 Homo sapiens similar to hypothetical protein 4932411E22 (LOC342600), mR
 XM_292627 Homo sapiens olfactory receptor, family 4, subfamily D, member 1 (OR4D1),

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XM_292664 Homo sapiens similar to NADH-ubiquinone oxidoreductase 15 kDa subunit ((
 XM_292674 Homo sapiens similar to leucine-rich repeat domain-containing protein (LOC:
 XM_292678 Homo sapiens similar to RIKEN cDNA 2610040E16 (LOC339291), mRNA
 XM_292700 Homo sapiens similar to 40S ribosomal protein S2 (LOC342808), mRNA
 XM_292707 Homo sapiens similar to TFIIF basal transcription factor complex p62 subunit
 XM_292717 Homo sapiens similar to KIAA1074 protein (LOC342850), mRNA
 XM_292723 Homo sapiens similar to zinc finger protein 495 (LOC342933), mRNA
 XM_292724 Homo sapiens similar to zinc finger protein 495 (LOC342934), mRNA
 XM_292729 Homo sapiens similar to double homeobox protein (LOC342939), mRNA
 XM_292740 Homo sapiens hypothetical protein LOC342892 (LOC342892), mRNA
 XM_292745 Homo sapiens similar to F-box only protein 2 (LOC342897), mRNA
 XM_292765 Homo sapiens zinc finger protein 404 (ZNF404), mRNA
 XM_292778 Homo sapiens leucine-rich repeats and immunoglobulin-like domains 4 (LRIC
 XM_292779 Homo sapiens hypothetical LOC342918 (LOC342918), mRNA
 XM_292784 Homo sapiens similar to hypothetical protein (LOC339351), mRNA
 XM_292785 Homo sapiens similar to contains transmembrane (TM) region (LOC342865),
 XM_292796 Homo sapiens similar to ret finger protein-like 1 (LOC342931), mRNA
 XM_292803 Homo sapiens similar to Hypothetical protein CBG18249 (LOC342958), mRNA
 XM_292810 Homo sapiens similar to Zinc finger protein 20 (Zinc finger protein KOK13) (C
 XM_292813 Homo sapiens similar to hypothetical protein FLJ38281 (LOC342972), mRNA
 XM_292817 Homo sapiens similar to actin 3 - fruit fly (*Drosophila melanogaster*) (fragment
 XM_292819 Homo sapiens nanos homolog 3 (*Drosophila*) (NANOS3), mRNA
 XM_292820 Homo sapiens hypothetical LOC342979 (LOC342979), mRNA
 XM_292824 Homo sapiens similar to hypothetical protein MGC45408 (LOC342969), mRNA
 XM_292832 Homo sapiens similar to hypothetical protein FLJ12488 (LOC342991), mRNA
 XM_292836 Homo sapiens similar to ribosomal protein L34; 60S ribosomal protein L34 (L
 XM_292850 Homo sapiens similar to pMesogenin1 (LOC343930), mRNA
 XM_292873 Homo sapiens similar to 201030C02Rik protein (LOC343990), mRNA
 XM_292889 Homo sapiens similar to Gnot1 homeodomain protein (LOC344022), mRNA
 XM_292895 Homo sapiens similar to zinc finger protein 135 (clone PHZ-17); zinc finger pr
 XM_292943 Homo sapiens Nck-associated protein 5 (NAP5), mRNA
 XM_292957 Homo sapiens similar to anaphase promoting complex subunit 1; anaphase-p
 XM_292958 Homo sapiens similar to Foxd1c protein (LOC344167), mRNA
 XM_292963 Homo sapiens similar to peptidylprolyl isomerase A (cyclophilin A) (LOC3441
 XM_292968 Homo sapiens similar to Homeobox even-skipped homolog protein 2 (EVX-2
 XM_292982 Homo sapiens similar to pote protein; Expressed in prostate, ovary, testis, ar
 XM_293018 Homo sapiens similar to Fatty acid-binding protein, epidermal (E-FABP) (Psc
 XM_293028 Homo sapiens similar to UNR-interacting protein (WD-40 repeat protein PT-V
 XM_293029 Homo sapiens similar to cyclin-dependent kinase-like 1 (CDC2-related kinase
 XM_293034 Homo sapiens similar to RIKEN cDNA 2010316F05 (LOC344405), mRNA
 XM_293042 Homo sapiens similar to ribosomal protein S12 (LOC344423), mRNA
 XM_293090 Homo sapiens FLJ00204 protein (FLJ00204), mRNA
 XM_293092 Homo sapiens G protein-coupled receptor 148 (GPR148), mRNA
 XM_293097 Homo sapiens similar to VsaA -like protein (LOC343565), mRNA
 XM_293104 Homo sapiens similar to dJ132F21.2 (Contains a novel protein similar to the
 XM_293106 Homo sapiens similar to hypothetical protein FLJ32191 (LOC343593), mRNA
 XM_293121 Homo sapiens similar to ba379F14.2 (novel protein) (LOC343629), mRNA
 XM_293123 Homo sapiens similar to dJ1100H13.4 (putative RhoGAP domain containing
 XM_293157 Homo sapiens similar to beta-galactoside alpha-2,3-sialyltransferase (LOC34
 XM_293160 Homo sapiens similar to dJ310O13.4 (novel protein similar to predicted C. el
 XM_293177 Homo sapiens similar to Zinc finger protein 85 (Zinc finger protein HPF4) (H1
 XM_293225 Homo sapiens similar to RIKEN cDNA 4930583C14 (LOC343854), mRNA
 XM_293226 Homo sapiens similar to POM121 membrane glycoprotein-like 1 (LOC343851)
 XM_293276 Homo sapiens similar to Heat shock 27 kDa protein (HSP 27) (Stress-respon
 XM_293284 Homo sapiens similar to Zinc finger protein 81 (HFZ20) (LOC347344), mRNA
 XM_293293 Homo sapiens similar to BMP-2 inducible protein kinase (BIKe) (HRIHFB201
 XM_293312 Homo sapiens similar to H3 histone, family 3B (LOC347376), mRNA

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XM_293320 Homo sapiens similar to bA370B6.1 (similar to histone H2B) (LOC347393), n
 XM_293325 Homo sapiens similar to ENSANGP00000015797 (LOC347411), mRNA
 XM_293332 Homo sapiens similar to RIKEN cDNA 1700113O17 (LOC340549), mRNA
 XM_293334 Homo sapiens similar to KIAA1726 protein (LOC340554), mRNA
 XM_293342 Homo sapiens similar to TGF beta-inducible nuclear protein 1 (Hairy cell leu
 XM_293352 Homo sapiens similar to P38IP protein (LOC347438), mRNA
 XM_293354 Homo sapiens similar to H326 (LOC347442), mRNA
 XM_293360 Homo sapiens similar to hSIAH1 (LOC340571), mRNA
 XM_293366 Homo sapiens similar to Olfactory receptor 13H1 (LOC347468), mRNA
 XM_293380 Homo sapiens similar to hypothetical protein A630014H24 (LOC347454), mF
 XM_293387 Homo sapiens similar to KIAA1892-like (LOC340578), mRNA
 XM_293396 Homo sapiens similar to Heat shock transcription factor, Y-linked (Heat shod
 XM_293398 Homo sapiens RAB41, member RAS homolog family (RAB41), mRNA
 XM_293401 Homo sapiens similar to arylsulfatase (LOC347527), mRNA
 XM_293405 Homo sapiens similar to hypothetical protein D430021I08 (LOC340595), mR
 XM_293407 Homo sapiens similar to melanoma antigen 2 (LOC347541), mRNA
 XM_293412 Homo sapiens similar to ribosomal protein L18a; 60S ribosomal protein L18a
 XM_293416 Homo sapiens similar to nuclear protein p30 (LOC347549), mRNA
 XM_293449 Homo sapiens similar to Cyorf16 protein (LOC389915), mRNA
 XM_293460 Homo sapiens similar to testis specific protein, Y-linked (LOC347596), mRN/
 XM_293514 Homo sapiens similar to CG17293-PA (LOC344620), mRNA
 XM_293529 Homo sapiens similar to RIKEN cDNA 4930558O21 (LOC344657), mRNA
 XM_293542 Homo sapiens similar to Elongation factor 1-gamma (EF-1-gamma) (eEF-1B
 XM_293565 Homo sapiens similar to seven transmembrane helix receptor (LOC344729),
 XM_293570 Homo sapiens similar to Heterogeneous nuclear ribonucleoprotein A1 (Helix-
 XM_293577 Homo sapiens similar to Arylacetamide deacetylase (AADAC) (LOC344752),
 XM_293580 Homo sapiens G protein-coupled receptor 149 (GPR149), mRNA
 XM_293581 Homo sapiens similar to seven transmembrane helix receptor (LOC344760),
 XM_293596 Homo sapiens similar to peptidylprolyl isomerase A (LOC344797), mRNA
 XM_293599 Homo sapiens type II transmembrane serine protease 7 (TMPRSS7), mRNA
 XM_293633 Homo sapiens similar to Protein MGC35450/QtsA-16602 (LOC344892), mR/
 XM_293656 Homo sapiens similar to ENSANGP00000007226 (LOC339951), mRNA
 XM_293669 Homo sapiens similar to actinin, alpha 4 (LOC344978), mRNA
 XM_293671 Homo sapiens similar to GTP-binding protein SAR1a (COPII-associated sma
 XM_293680 Homo sapiens similar to MGC53273 protein (LOC345051), mRNA
 XM_293687 Homo sapiens similar to RIKEN cDNA 5730467H21 (LOC345079), mRNA
 XM_293715 Homo sapiens similar to CG13722-PA (LOC345158), mRNA
 XM_293745 Homo sapiens hypothetical LOC345222 (LOC345222), mRNA
 XM_293801 Homo sapiens similar to CG32656-PA (LOC345375), mRNA
 XM_293802 Homo sapiens similar to UDP-glucuronosyltransferase 2B15 precursor, micr
 XM_293821 Homo sapiens similar to profilin 3 (LOC345456), mRNA
 XM_293828 Homo sapiens similar to hypothetical protein 9630041N07 (LOC345462), mF
 XM_293829 Homo sapiens similar to UNR protein (LOC345463), mRNA
 XM_293868 Homo sapiens similar to hypothetical protein (LOC345537), mRNA
 XM_293875 Homo sapiens similar to RIKEN cDNA B130016O10 gene (LOC345557), mR
 XM_293886 Homo sapiens similar to Ubiquitin carboxyl-terminal hydrolase 7 (Ubiquitin th
 XM_293893 Homo sapiens LRG-47-like protein (LRG47), mRNA
 XM_293903 Homo sapiens similar to fibrillarin (LOC345630), mRNA
 XM_293911 Homo sapiens hypothetical protein FLJ40191 (FLJ40191), mRNA
 XM_293918 Homo sapiens similar to geminin (LOC345643), mRNA
 XM_293923 Homo sapiens similar to protease (prosome, macropain) 26S subunit, ATPas
 XM_293924 Homo sapiens similar to RIKEN cDNA 4732495G21 gene (LOC345651), mR
 XM_293927 Homo sapiens similar to template acylating factor-I alpha (LOC345659), mF
 XM_293937 Homo sapiens similar to RIKEN cDNA 0610012A05 (LOC345711), mRNA
 XM_293943 Homo sapiens similar to histone (15.4 kD) (his-72) (LOC340096), mRNA
 XM_293971 Homo sapiens similar to hypothetical protein (LOC345778), mRNA
 XM_293976 Homo sapiens similar to RIKEN cDNA 6430502M16 gene (LOC340120), mR

XM_293984 Homo sapiens similar to Transcription factor BTF3 (RNA polymerase B trans
 XM_294017 Homo sapiens solute carrier family 35, member D3 (SLC35D3), mRNA
 XM_294019 Homo sapiens similar to Ect2 protein (LOC345930), mRNA
 XM_294070 Homo sapiens similar to Glyceraldehyde 3-phosphate dehydrogenase, liver (L
 XM_294077 Homo sapiens similar to dJ153G14.3 (novel C2H2 type Zinc Finger protein) (L
 XM_294093 Homo sapiens similar to bA145L22.2 (novel KRAB box containing C2H2 type
 XM_294139 Homo sapiens chromosome 6 open reading frame 143 (C6orf143), mRNA
 XM_294165 Homo sapiens similar to septin 10 isoform 1 (LOC346288), mRNA
 XM_294209 Homo sapiens similar to Unc4.1 homeobox (LOC340260), mRNA
 XM_294219 Homo sapiens similar to RIKEN cDNA A930017N06 gene (LOC346355), mR
 XM_294247 Homo sapiens similar to Splicing factor, arginine/serine-rich, 46kD (LOC3484
 XM_294249 Homo sapiens similar to GluR-delta2 philic-protein (LOC340265), mRNA
 XM_294261 Homo sapiens similar to hypothetical protein FLJ22527 (LOC346545), mRN/
 XM_294265 Homo sapiens similar to envelope protein (LOC346547), mRNA
 XM_294310 Homo sapiens similar to Olfactory receptor 6V1 (LOC346517), mRNA
 XM_294311 Homo sapiens similar to Histidine triad nucleotide-binding protein 1 (Adenosl
 XM_294366 Homo sapiens similar to Olfactory receptor 2A12 (LOC346525), mRNA
 XM_294318 Homo sapiens similar to Olfactory receptor 2A1 (LOC346528), mRNA
 XM_294319 Homo sapiens similar to Importin alpha-2 subunit (Karyopherin alpha-2 subu
 XM_294328 Homo sapiens similar to Protein C6orf66 (HSPC125) (My013 protein) (LOC3
 XM_294353 Homo sapiens similar to RIKEN cDNA 6332401019 gene (LOC340344), mR
 XM_294354 Homo sapiens similar to 40S ribosomal protein S2 (LOC346643), mRNA
 XM_294357 Homo sapiens hypothetical protein LOC346653 (LOC346653), mRNA
 XM_294395 Homo sapiens similar to Na+/L-ascorbic acid transporter 1; SVCT1 (LOC346
 XM_294370 Homo sapiens guanine nucleotide binding protein, alpha transducing 3 (GNA
 XM_294383 Homo sapiens similar to seven transmembrane helix receptor (LOC346708),
 XM_294387 Homo sapiens similar to hypothetical protein 8230402K04 (LOC340359), mR
 XM_294438 Homo sapiens similar to 60S RIBOSOMAL PROTEIN L5 (LOC346848), mR
 XM_294450 Homo sapiens similar to solute carrier family 16 (monocarboxylic acid transp
 XM_294456 Homo sapiens similar to putative nuclear protein (4B256) (LOC346910), mR
 XM_294468 Homo sapiens similar to H2A histone family, member Z (LOC346990), mRN/
 XM_294473 Homo sapiens similar to ribosomal protein L37 (LOC346950), mRNA
 XM_294476 Homo sapiens similar to Interferon-induced transmembrane protein 3 (Interfe
 XM_294480 Homo sapiens similar to pote protein; Expressed in prostate, ovary, testis, ar
 XM_294521 Homo sapiens FLJ43950 protein (FLJ43950), mRNA
 XM_294531 Homo sapiens similar to Olfactory receptor 1J1 (LOC347168), mRNA
 XM_294533 Homo sapiens olfactory receptor, family 1, subfamily J, member 4 (OR1J4), r
 XM_294534 Homo sapiens similar to Olfactory receptor 1B1 (Olfactory receptor 9-B) (OR
 XM_294540 Homo sapiens similar to cancer related gene-liver 1 (LOC340485), mRNA
 XM_294567 Homo sapiens similar to bA113O24.1 (similar to insulin-like growth factor bin
 XM_294568 Homo sapiens similar to transcription elongation factor IIS - mouse (LOC340
 XM_294574 Homo sapiens similar to A-kinase anchor protein 8; A-kinase anchor protein,
 XM_294581 Homo sapiens similar to ribosomal protein L36; 60S ribosomal protein L36 (L
 XM_294584 Homo sapiens similar to hypothetical protein FLJ40432 (LOC347262), mRN/
 XM_294590 Homo sapiens similar to bA13B9.3 (novel protein similar to KRT8) (LOC3472
 XM_294592 Homo sapiens similar to RIKEN cDNA 2310039E09 (LOC347273), mRNA
 XM_294634 Homo sapiens similar to alpha-2 macroglobulin family protein VIP (LOC3402
 XM_294686 Homo sapiens hypothetical LOC338616 (LOC338616), mRNA
 XM_294675 Homo sapiens hypothetical protein LOC338667 (LOC338667), mRNA
 XM_294680 Homo sapiens hypothetical protein LOC338694 (LOC338694), mRNA
 XM_294688 Homo sapiens hypothetical LOC338731 (LOC338731), mRNA
 XM_294692 Homo sapiens hypothetical LOC338749 (LOC338749), mRNA
 XM_294723 Homo sapiens hypothetical LOC338825 (LOC338825), mRNA
 XM_294743 Homo sapiens hypothetical LOC338918 (LOC338918), mRNA
 XM_294750 Homo sapiens hypothetical LOC338951 (LOC338951), mRNA
 XM_294751 Homo sapiens similar to RIKEN cDNA 4930425N13 (LOC338949), mRNA
 XM_294785 Homo sapiens similar to FLJ40113 protein (LOC388154), mRNA

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XM_294775 Homo sapiens hypothetical LOC339022 (LOC339022), mRNA
 XM_294778 Homo sapiens hypothetical LOC339025 (LOC339025), mRNA
 XM_294794 Homo sapiens similar to putative membrane-bound dipeptidase-2 (LOC3390)
 XM_294802 Homo sapiens hypothetical LOC339077 (LOC339077), mRNA
 XM_294854 Homo sapiens hypothetical LOC339209 (LOC339209), mRNA
 XM_294867 Homo sapiens hypothetical LOC339226 (LOC339226), mRNA
 XM_294894 Homo sapiens hypothetical LOC339281 (LOC339281), mRNA
 XM_294906 Homo sapiens hypothetical LOC339306 (LOC339306), mRNA
 XM_294910 Homo sapiens hypothetical LOC339352 (LOC339352), mRNA
 XM_294914 Homo sapiens hypothetical LOC339358 (LOC339358), mRNA
 XM_294919 Homo sapiens hypothetical protein LOC339366 (LOC339366), mRNA
 XM_294922 Homo sapiens hypothetical LOC339375 (LOC339375), mRNA
 XM_294960 Homo sapiens hypothetical LOC339453 (LOC339453), mRNA
 XM_294963 Homo sapiens LOC388583 (LOC388583), mRNA
 XM_294993 Homo sapiens hypothetical protein LOC339529 (LOC339529), mRNA
 XM_294997 Homo sapiens hypothetical LOC339541 (LOC339541), mRNA
 XM_295007 Homo sapiens hypothetical LOC339583 (LOC339583), mRNA
 XM_295017 Homo sapiens chromosome 21 open reading frame 54 (C21orf54), mRNA
 XM_295034 Homo sapiens hypothetical LOC339693 (LOC339693), mRNA
 XM_295058 Homo sapiens hypothetical LOC339760 (LOC339760), mRNA
 XM_295059 Homo sapiens hypothetical LOC339771 (LOC339771), mRNA
 XM_295062 Homo sapiens hypothetical LOC339782 (LOC339782), mRNA
 XM_295061 Homo sapiens hypothetical LOC339875 (LOC339875), mRNA
 XM_295066 Homo sapiens hypothetical LOC339899 (LOC339899), mRNA
 XM_295097 Homo sapiens hypothetical LOC339902 (LOC339902), mRNA
 XM_295120 Homo sapiens hypothetical LOC339985 (LOC339985), mRNA
 XM_295126 Homo sapiens hypothetical LOC339997 (LOC339997), mRNA
 XM_295146 Homo sapiens hypothetical LOC340065 (LOC340065), mRNA
 XM_295155 Homo sapiens hypothetical protein LOC340094 (LOC340094), mRNA
 XM_295166 Homo sapiens hypothetical LOC340148 (LOC340148), mRNA
 XM_295169 Homo sapiens hypothetical LOC340149 (LOC340149), mRNA
 XM_295178 Homo sapiens hypothetical LOC340171 (LOC340171), mRNA
 XM_295195 Homo sapiens similar to Matn2-prov protein (LOC340267), mRNA
 XM_295200 Homo sapiens hypothetical LOC340286 (LOC340286), mRNA
 XM_295205 Homo sapiens hypothetical LOC340349 (LOC340349), mRNA
 XM_295213 Homo sapiens hypothetical LOC340346 (LOC340346), mRNA
 XM_295252 Homo sapiens hypothetical LOC340450 (LOC340450), mRNA
 XM_295257 Homo sapiens hypothetical LOC340477 (LOC340477), mRNA
 XM_295261 Homo sapiens hypothetical LOC340511 (LOC340511), mRNA
 XM_295263 Homo sapiens hypothetical protein LOC340508 (LOC340508), mRNA
 XM_295270 Homo sapiens hypothetical protein LOC340581 (LOC340581), mRNA
 XM_295309 Homo sapiens similar to putative neuronal cell adhesion molecule (LOC3430)
 XM_295598 Homo sapiens hypothetical LOC343484 (LOC343484), mRNA
 XM_295865 Homo sapiens similar to apical early endosomal glycoprotein precursor (LOC
 XM_296117 Homo sapiens similar to Heterogeneous nuclear ribonucleoprotein A1 (Helix-
 XM_296315 Homo sapiens similar to 60S ribosomal protein L35 (LOC341604), mRNA
 XM_296817 Homo sapiens similar to hypothetical protein DKFZp434P0316 (LOC342346)
 XM_297205 Homo sapiens hypothetical LOC342900 (LOC342900), mRNA
 XM_297687 Homo sapiens similar to transmembrane protein 16C; chromosome 11 open
 XM_297816 Homo sapiens similar to dJ824F16.3 (novel protein similar to mouse thromb
 XM_298045 Homo sapiens hypothetical LOC347475 (LOC347475), mRNA
 XM_298053 Homo sapiens hypothetical LOC347487 (LOC347487), mRNA
 XM_298151 Homo sapiens hypothetical LOC344595 (LOC344595), mRNA
 XM_298233 Homo sapiens similar to hypothetical protein FLJ38348 (LOC344709), mRNA
 XM_301210 Homo sapiens similar to testis-specific protein NYD-TSP1 (LOC388775), mR
 XM_350780 Homo sapiens KIAA0592 protein (KIAA0592), mRNA
 XM_350880 Homo sapiens ras homolog gene family, member C like 1 (ARHCL1), mRNA

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XM_351335 Homo sapiens similar to hypothetical protein KIAA0454 - human (fragment) (

XM_351366 Homo sapiens similar to dJ309K20.1.1 (novel protein similar to dysferlin, iso

XM_351473 Homo sapiens similar to GLYCEROL-3-PHOSPHATE ACYLTRANSFERASE

XM_351574 Homo sapiens similar to Succinate dehydrogenase [ubiquinone] flavoprotein

XM_351617 Homo sapiens similar to dJ28I24.1.2 (Spinal Muscular Atrophy region (SMA3

XM_351723 Homo sapiens similar to tripartite motif protein TRIM50B (LOC375593), mRNA

XM_351854 Homo sapiens similar to cDNA sequence BC034076 (LOC389797), mRNA

XM_351948 Homo sapiens similar to Transcript Y 7 protein (LOC389916), mRNA

XM_352159 Homo sapiens similar to hypothetical protein LOC339047 (LOC388215), mR

XM_352463 Homo sapiens similar to myoferlin isoform b; fer-1-like 3 (LOC391408), mRNA

XM_352847 Homo sapiens hypothetical protein LOC340529 (LOC340529), mRNA

XM_353628 Homo sapiens similar to tripartite motif protein TRIM50B (LOC378125), mRNA

XM_370632 Homo sapiens similar to ubiquitin-conjugating enzyme E2 variant 1 isoform a

XM_370533 Homo sapiens similar to eukaryotic initiation factor 4B (LOC387637), mRNA

XM_370534 Homo sapiens similar to RIKEN cDNA E130012A19 (LOC387640), mRNA

XM_370536 Homo sapiens similar to RIKEN cDNA 4921622E24 (LOC387642), mRNA

XM_370537 Homo sapiens similar to amyloid beta (A4) precursor protein-binding, family I

XM_370538 Homo sapiens similar to amyloid beta (A4) precursor protein-binding, family I

XM_370541 Homo sapiens FLJ44037 protein (FLJ44037), mRNA

XM_370542 Homo sapiens similar to supervillin isoform 2; membrane-associated F-actin

XM_370543 Homo sapiens similar to Pre-B cell enhancing factor precursor (LOC387651)

XM_370545 Homo sapiens similar to bA291L22.2 (similar to CDC10 (cell division cycle 1)

XM_370554 Homo sapiens similar to ARF GTPase-activating protein (LOC387671), mRNA

XM_370555 Homo sapiens similar to Glutamate dehydrogenase, mitochondrial precursor

XM_370557 Homo sapiens similar to KIAA0592 protein (LOC387676), mRNA

XM_370560 Homo sapiens similar to ADP-ribosylation factor-like protein 4 (LOC387684),

XM_370562 Homo sapiens LOC399782 (LOC387688), mRNA

XM_370564 Homo sapiens similar to BMS1-like, ribosome assembly protein; ribosome bi

XM_370565 Homo sapiens similar to hypothetical protein FLJ20967 (LOC387694), mRNA

XM_370566 Homo sapiens similar to RLLV1833 (LOC387695), mRNA

XM_370567 Homo sapiens KIAA1975 protein similar to MRIP2 (KIAA1975), mRNA

XM_370569 Homo sapiens similar to expressed sequence AW210596 (LOC387700), mRNA

XM_370570 Homo sapiens similar to hypothetical protein FLJ25224 (LOC387702), mRNA

XM_370571 Homo sapiens similar to ATP-dependent DNA helicase II, 70 kDa subunit (Lu

XM_370573 Homo sapiens similar to KIAA1345 protein (LOC387707), mRNA

XM_370575 Homo sapiens hypothetical protein MGC11279 (MGC11279), mRNA

XM_370577 Homo sapiens similar to RIKEN cDNA 6430537H07 gene (LOC387712), mRNA

XM_370580 Homo sapiens similar to Hmx2 protein (LOC387716), mRNA

XM_370582 Homo sapiens hypothetical gene supported by BC062717 (LOC387718), mRNA

XM_370584 Homo sapiens hypothetical gene supported by AK127642 (LOC387720), mRNA

XM_370586 Homo sapiens similar to bA442O18.2 (novel protein) (LOC387721), mRNA

XM_370588 Homo sapiens LOC399826 (LOC387723), mRNA

XM_370589 Homo sapiens similar to double homeobox protein (LOC387727), mRNA

XM_370591 Homo sapiens similar to hypothetical protein MGC5560 (LOC387728), mRNA

XM_370593 Homo sapiens similar to double homeobox protein (LOC387729), mRNA

XM_370597 Homo sapiens similar to putative haemopoietic membrane protein (LOC3877

XM_370603 Homo sapiens similar to SB153 protein (LOC387745), mRNA

XM_370605 Homo sapiens similar to seven transmembrane helix receptor (LOC387748),

XM_370606 Homo sapiens hypothetical gene supported by BX647806 (LOC387749), mRNA

XM_370607 Homo sapiens similar to SlzC220F6.1 (novel protein similar to human dynei

XM_370610 Homo sapiens similar to 60S ribosomal protein L23a (LOC387752), mRNA

XM_370611 Homo sapiens similar to 60S ribosomal protein L21 (LOC387753), mRNA

XM_370612 Homo sapiens calcitonin-related polypeptide, beta (CALCB), mRNA

XM_370613 Homo sapiens similar to RIKEN cDNA 3830422K02 (LOC387755), mRNA

XM_370615 Homo sapiens hypothetical protein DKFPz686O24166 (DKFPz686O24166),

XM_370616 Homo sapiens hypothetical protein LOC338645 (LOC338645), mRNA

XM_370618 Homo sapiens hypothetical protein FLJ20294 (FLJ20294), mRNA

XM_370619 Homo sapiens similar to mitochondrial carrier protein MGC4399 (LOC38776; XM_370621 Homo sapiens similar to tripartite motif protein 48; TRIM48 (LOC387770), mF
XM_370622 Homo sapiens similar to seven transmembrane helix receptor (LOC387772), mF
XM_370623 Homo sapiens similar to MGC15937 protein (LOC387773), mRNA
XM_370629 Homo sapiens similar to organic anion transporter 6 (LOC387775), mRNA
XM_370630 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 14B (PPF
XM_370631 Homo sapiens similar to p33 ringo (LOC387778), mRNA
XM_370632 Homo sapiens similar to phosphoglycerate mutase 1 (brain) (LOC387779), mF
XM_370633 Homo sapiens similar to solute carrier family 22 member 6; organic cationic 1
XM_370634 Homo sapiens similar to UBIQUITIN-LIKE PROTEIN FUBI (LOC387781), mF
XM_370635 Homo sapiens KIAA0280 protein (KIAA0280), mRNA
XM_370636 Homo sapiens similar to RIKEN cDNA 2610209A20 (LOC387787), mRNA
XM_370638 Homo sapiens LOC399929 (LOC387790), mRNA
XM_370639 Homo sapiens similar to Mitochondrial import receptor subunit TOM20 homol
XM_370642 Homo sapiens LOC399932 (LOC387795), mRNA
XM_370644 Homo sapiens similar to tripartite motif-containing 43 (LOC387800), mRNA
XM_370648 Homo sapiens LOC399943 (LOC387804), mRNA
XM_370649 Homo sapiens similar to ARP2/3 complex 21 kDa subunit (p21-ARC) (Actin-r
XM_370651 Homo sapiens hypothetical protein FLJ32810 (FLJ32810), mRNA
XM_370652 Homo sapiens dynein, cytoplasmic, heavy polypeptide 2 (DNCH2), mRNA
XM_370653 Homo sapiens KIAA1826 protein (KIAA1826), mRNA
XM_370654 Homo sapiens KIAA1726 protein (KIAA1726), mRNA
XM_370657 Homo sapiens placenta expressed transcript 1 (PLET1), mRNA
XM_370658 Homo sapiens similar to hypothetical protein FLJ25224 (LOC387811), mRNA
XM_370680 Homo sapiens KIAA1201 protein (KIAA1201), mRNA
XM_370681 Homo sapiens similar to seven transmembrane helix receptor (LOC387815),
XM_370682 Homo sapiens olfactory receptor, family 8, subfamily G, member 2 (OR8G2),
XM_370683 Homo sapiens retinoblastoma inhibiting gene 1 (RBIG1), mRNA
XM_370684 Homo sapiens similar to hypothetical protein (LOC387816), mRNA
XM_370685 Homo sapiens similar to DnaJ (Hsp40) homolog, subfamily B, member 6 isoform
XM_370686 Homo sapiens similar to hypothetical protein (LOC387822), mRNA
XM_370687 Homo sapiens KIAA1110 protein (KIAA1110), mRNA
XM_370688 Homo sapiens similar to ribosomal protein L13a; 60S ribosomal protein L13a
XM_370689 Homo sapiens homeobox C14 (LOC380030), mRNA
XM_370672 Homo sapiens similar to hypothetical protein SB153 isoform 1 (LOC387830),
XM_370674 Homo sapiens similar to helicase (LOC387832), mRNA
XM_370675 Homo sapiens similar to DKFZp434C0631 protein (LOC387834), mRNA
XM_370678 Homo sapiens similar to INFE5792 (LOC387836), mRNA
XM_370681 Homo sapiens similar to ribosomal protein L13a; 60S ribosomal protein L13a
XM_370682 Homo sapiens KIAA1467 protein (KIAA1467), mRNA
XM_370684 Homo sapiens similar to Elongation factor 1-alpha 1 (EF-1-alpha-1) (Elongat
XM_370685 Homo sapiens hypothetical protein LOC144363 (LOC144363), mRNA
XM_370686 Homo sapiens similar to RIKEN cDNA 2210417D09 (LOC387849), mRNA
XM_370687 Homo sapiens similar to ribosomal protein L13a; 60S ribosomal protein L13a
XM_370688 Homo sapiens similar to Adenylate kinase isoenzyme 4, mitochondrial (ATP-
XM_370690 Homo sapiens AT rich interactive domain 2 (ARID, RFX-like) (ARID2), mRNA
XM_370691 Homo sapiens similar to expressed sequence A1836003 (LOC387856), mRNA
XM_370692 Homo sapiens hypothetical protein LOC121006 (LOC121006), mRNA
XM_370693 Homo sapiens POU domain, class 6, transcription factor 1 (POU6F1), mRNA
XM_370695 Homo sapiens similar to K⁺ channel tetramerization protein (LOC387861), m
XM_370696 Homo sapiens hypothetical protein FLJ34236 (FLJ34236), mRNA
XM_370697 Homo sapiens similar to 40S ribosomal protein SA (P40) (34/67 kDa laminin
XM_370699 Homo sapiens protein tyrosine phosphatase, receptor type, Q (PTPRQ), mR
XM_370702 Homo sapiens FYVE, RhoGEF and PH domain containing 6 (FGD6), mRNA
XM_370704 Homo sapiens similar to 10 KD HEAT SHOCK PROTEIN, MITOCHONDRIAL
XM_370705 Homo sapiens LOC400067 (LOC387881), mRNA
XM_370707 Homo sapiens similar to hypothetical protein C130069F04 (LOC387890), mF

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XM_370709 Homo sapiens LOC400085 (LOC387894), mRNA
 XM_370710 Homo sapiens similar to protein 40kD (LOC387902), mRNA
 XM_370711 Homo sapiens similar to hypothetical protein (LOC387904), mRNA
 XM_370713 Homo sapiens similar to bA271B5.1 (similar to ribosomal protein S7) (LOC387901)
 XM_370714 Homo sapiens similar to Ferritin heavy chain (Ferritin H subunit) (LOC387901)
 XM_370715 Homo sapiens similar to hypothetical protein MGC48915 (LOC387911), mRNA
 XM_370716 Homo sapiens similar to TPTE and PTEN homologous inositol lipid phosphatase
 XM_370718 Homo sapiens similar to WGAR9166 (LOC387914), mRNA
 XM_370721 Homo sapiens similar to bA251J8.3.1 (novel protein, isoform 1) (LOC387920)
 XM_370722 Homo sapiens similar to RIKEN cDNA 8030451K01 (LOC387921), mRNA
 XM_370723 Homo sapiens similar to tubulin, beta 5 (LOC387922), mRNA
 XM_370724 Homo sapiens similar to ribosome associated membrane protein 4 (LOC387922), mRNA
 XM_370725 Homo sapiens similar to KIAA1612 protein (LOC387924), mRNA
 XM_370726 Homo sapiens similar to BB049667 protein (LOC387927), mRNA
 XM_370727 Homo sapiens similar to ribosomal protein L13a; 60S ribosomal protein L13a
 XM_370728 Homo sapiens similar to heterogeneous nuclear ribonucleoprotein A3 (LOC387928)
 XM_370729 Homo sapiens similar to Fatty acid-binding protein, epidermal (E-FABP) (Psc)
 XM_370731 Homo sapiens similar to expressed sequence AW121567 (LOC387944), mRNA
 XM_370732 Homo sapiens similar to expressed sequence AW121567 (LOC387945), mRNA
 XM_370733 Homo sapiens LOC400166 (LOC387949), mRNA
 XM_370734 Homo sapiens hypothetical gene supported by BC034570 (LOC387952), mRNA
 XM_370737 Homo sapiens hypothetical protein FLJ10357 (FLJ10357), mRNA
 XM_370738 Homo sapiens helicase with SNF2 domain 1 (HLSNF1), mRNA
 XM_370754 Homo sapiens thiamine triphosphatase (THTPA), mRNA
 XM_370756 Homo sapiens KIAA1305 (KIAA1305), mRNA
 XM_370758 Homo sapiens hypothetical gene supported by BX248251 (LOC387978), mRNA
 XM_370759 Homo sapiens similar to RIKEN cDNA D930036F22 gene (LOC387979), mRNA
 XM_370760 Homo sapiens similar to 60S RIBOSOMAL PROTEIN L12 (LOC387982), mRNA
 XM_370762 Homo sapiens similar to TIMM9 (LOC387990), mRNA
 XM_370763 Homo sapiens similar to ribosomal protein L31 (LOC387991), mRNA
 XM_370765 Homo sapiens papilin, proteoglycan-like sulfated glycoprotein (PAPLN), mRNA
 XM_370767 Homo sapiens chromosome 14 open reading frame 46 (C14orf46), mRNA
 XM_370768 Homo sapiens similar to Acylphosphatase, organ-common type isozyme (Ac)
 XM_370769 Homo sapiens hypothetical protein LOC161394 (LOC161394), mRNA
 XM_370772 Homo sapiens similar to protease inhibitor (LOC388007), mRNA
 XM_370776 Homo sapiens similar to RT11 (LOC388015), mRNA
 XM_370777 Homo sapiens similar to Lysophospholipase (LOC374569), mRNA
 XM_370778 Homo sapiens similar to expressed sequence AJ839735 (LOC388021), mRNA
 XM_370779 Homo sapiens hypothetical gene supported by AK131040 (LOC388022), mRNA
 XM_370781 Homo sapiens similar to Ig alpha-2 chain C region (LOC388025), mRNA
 XM_370782 Homo sapiens similar to Ig epsilon chain C region (LOC388026), mRNA
 XM_370785 Homo sapiens chromosome 14 open reading frame 81 (C14orf81), mRNA
 XM_370826 Homo sapiens similar to breast cancer antigen NY-BR-1 (LOC388065), mRNA
 XM_370829 Homo sapiens similar to carnitine deficiency-associated gene expressed in v
 XM_370830 Homo sapiens similar to breast cancer anti-estrogen resistance 1; Crk-associ
 XM_370831 Homo sapiens similar to Ribosome biogenesis protein BMS1 homolog (LOC388065)
 XM_370832 Homo sapiens similar to Ribosome biogenesis protein BMS1 homolog (LOC388065)
 XM_370833 Homo sapiens similar to 40S ribosomal protein S8 (LOC388076), mRNA
 XM_370834 Homo sapiens similar to immunoglobulin heavy chain variable region (LOC388076)
 XM_370835 Homo sapiens similar to FLJ27099 protein (LOC388078), mRNA
 XM_370836 Homo sapiens similar to ZCCHC2 protein (LOC388079), mRNA
 XM_370837 Homo sapiens similar to hypothetical protein (LOC388080), mRNA
 XM_370838 Homo sapiens hypothetical protein LOC339005 (LOC339005), mRNA
 XM_370839 Homo sapiens similar to golgin-67 isoform b (LOC388084), mRNA
 XM_370840 Homo sapiens similar to hypothetical protein (LOC388085), mRNA
 XM_370843 Homo sapiens similar to hypothetical protein (LOC388092), mRNA
 XM_370844 Homo sapiens similar to hypothetical protein (LOC388094), mRNA

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XM_370845 Homo sapiens similar to hyperpolarization activated cyclic nucleotide-gated γ
 XM_370846 Homo sapiens similar to hypothetical protein FLJ35785 (LOC388098), mRNA/
 XM_370848 Homo sapiens similar to chromosome 1 open reading frame 37 (LOC388104
 XM_370849 Homo sapiens similar to hyperpolarization activated cyclic nucleotide-gated γ
 XM_370851 Homo sapiens similar to hypothetical protein FLJ35785 (LOC388109), mRNA/
 XM_370852 Homo sapiens similar to 4930563P21Rik protein (LOC388110), mRNA
 XM_370853 Homo sapiens similar to Nanog homeobox; homeobox transcription factor Nr
 XM_370855 Homo sapiens similar to RIKEN cDNA 2600011L02 (LOC388115), mRNA
 XM_370856 Homo sapiens similar to RIKEN cDNA 6720467C03 (LOC388116), mRNA
 XM_370858 Homo sapiens similar to kinesin-like protein (LOC388118), mRNA
 XM_370863 Homo sapiens ATPase, Class I, type 8B, member 4 (ATP8B4), mRNA
 XM_370864 Homo sapiens similar to TNF-induced protein (LOC388121), mRNA
 XM_370865 Homo sapiens similar to Laminin receptor 1 (LOC388122), mRNA
 XM_370866 Homo sapiens hypothetical protein FLJ25756 (FLJ25756), mRNA
 XM_370867 Homo sapiens hypothetical protein FLJ20086 (FLJ20086), mRNA
 XM_370868 Homo sapiens similar to hypothetical LOC237397 (LOC388125), mRNA
 XM_370871 Homo sapiens hypothetical protein LOC145837 (LOC145837), mRNA
 XM_370872 Homo sapiens similar to 60S ribosomal protein L17 (L23) (LOC388132), mRi
 XM_370873 Homo sapiens similar to RIKEN cDNA 6030419C18 gene (LOC388135), mR
 XM_370876 Homo sapiens similar to Golgi autoantigen, golgin subfamily A member 6 (Gc
 XM_370878 Homo sapiens KIAA2002 protein (KIAA2002), mRNA
 XM_370879 Homo sapiens similar to 60S ribosomal protein L21 (LOC388143), mRNA
 XM_370880 Homo sapiens mesoderm development candidate 2 (MESDC2), mRNA
 XM_370881 Homo sapiens similar to FLJ40113 protein (LOC388146), mRNA
 XM_370882 Homo sapiens similar to ribosomal protein L9; 60S ribosomal protein L9 (Loi
 XM_370883 Homo sapiens LOC400417 (LOC388148), mRNA
 XM_370886 Homo sapiens similar to FLJ40113 protein (LOC388151), mRNA
 XM_370887 Homo sapiens LOC400422 (LOC388153), mRNA
 XM_370893 Homo sapiens similar to hypothetical protein FLJ13119 (LOC388159), mRNA/
 XM_370894 Homo sapiens LOC400427 (LOC388160), mRNA
 XM_370895 Homo sapiens LOC400432 (LOC388163), mRNA
 XM_370897 Homo sapiens similar to RIKEN cDNA 3010021M21 (LOC388165), mRNA
 XM_370898 Homo sapiens similar to FLJ40113 protein (LOC388166), mRNA
 XM_370899 Homo sapiens similar to FLJ40113 protein (LOC388167), mRNA
 XM_370904 Homo sapiens similar to hypothetical protein 4921538O11 (LOC388173), mF
 XM_370905 Homo sapiens similar to Golgi autoantigen, golgin subfamily a, 2; golgin-95; i
 XM_370906 Homo sapiens similar to hypothetical protein FLJ20147 (LOC388175), mRNA/
 XM_370908 Homo sapiens similar to cyclin-E binding protein 1 (H. sapiens) (MGC14386)
 XM_370909 Homo sapiens similar to H2A histone family, member V isoform 2; histone H;
 XM_370910 Homo sapiens similar to KIAA0974 protein (LOC388181), mRNA
 XM_370911 Homo sapiens hypothetical gene supported by AK124283 (LOC388182), mR
 XM_370917 Homo sapiens similar to hypothetical protein (LOC388189), mRNA
 XM_370918 Homo sapiens hypothetical protein DKFZp434P162 (DKFZp434P162), mRN
 XM_370924 Homo sapiens LOC400486 (LOC388199), mRNA
 XM_370925 Homo sapiens hypothetical protein LOC283951 (LOC283951), mRNA
 XM_370926 Homo sapiens similar to KIAA0445 protein (LOC388202), mRNA
 XM_370927 Homo sapiens ring finger protein 151 (RNF151), mRNA
 XM_370928 Homo sapiens KIAA1171 protein (KIAA1171), mRNA
 XM_370930 Homo sapiens similar to RIKEN cDNA 1520401A03 gene (LOC388205), mRi
 XM_370931 Homo sapiens olfactory receptor, family 1, subfamily F, member 2 (OR1F2),
 XM_370932 Homo sapiens hypothetical protein FLJ39639 (FLJ39639), mRNA
 XM_370934 Homo sapiens similar to CG15828-PA (LOC388210), mRNA
 XM_370935 Homo sapiens LOC400498 (LOC388211), mRNA
 XM_370938 Homo sapiens similar to QRWTS810 (LOC388218), mRNA
 XM_370939 Homo sapiens similar to KIAA0220 (LOC388221), mRNA
 XM_370942 Homo sapiens similar to carbonic anhydrase VA, mitochondrial precursor; ca
 XM_370943 Homo sapiens similar to MGC9515 protein (LOC400510), mRNA

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XM_370944 Homo sapiens hypothetical protein LOC146177 (LOC146177), mRNA
 XM_370946 Homo sapiens similar to protein kinase/ribonuclease IRE1 beta (LOC388226)
 XM_370947 Homo sapiens ER to nucleus signalling 2 (ERN2), mRNA
 XM_370948 Homo sapiens similar to SH3-binding kinase (LOC388228), mRNA
 XM_370949 Homo sapiens similar to Group X secretory phospholipase A2 precursor (Phx
 XM_370952 Homo sapiens similar to MGC9515 protein (LOC388233), mRNA
 XM_370958 Homo sapiens similar to nuclear pore complex interacting protein (LOC3882
 XM_370959 Homo sapiens similar to RRN3 (LOC388238), mRNA
 XM_370965 Homo sapiens similar to hypothetical protein BCD11981 (LOC388242), mRNA
 XM_370966 Homo sapiens similar to carbonic anhydrase VA, mitochondrial precursor; ca
 XM_370967 Homo sapiens hypothetical protein LOC124411 (LOC124411), mRNA
 XM_370968 Homo sapiens similar to KIAA1501 protein (LOC388248), mRNA
 XM_370972 Homo sapiens similar to Adrenoleukodystrophy protein (ALDP) (LOC388253
 XM_370973 Homo sapiens similar to KIAA1501 protein (LOC388255), mRNA
 XM_370974 Homo sapiens similar to NY-REN-7 antigen (LOC388258), mRNA
 XM_370975 Homo sapiens similar to protein kinase related to Raf protein kinases; Metho
 XM_370977 Homo sapiens similar to Immunoglobulin heavy chain VH3 (LOC388264), mF
 XM_370980 Homo sapiens similar to RAB41 (LOC388271), mRNA
 XM_370981 Homo sapiens similar to RIKEN cDNA 4921524J17 (LOC388272), mRNA
 XM_370982 Homo sapiens similar to Heterogeneous nuclear ribonucleoprotein A1 (Helix-
 XM_370984 Homo sapiens LOC388284 (LOC388284), mRNA
 XM_370985 Homo sapiens similar to hypothetical protein (L1H 3 region) - human (LOC3E
 XM_370987 Homo sapiens LOC388289 (LOC388289), mRNA
 XM_370988 Homo sapiens similar to protein 40kD (LOC388290), mRNA
 XM_370991 Homo sapiens similar to hypothetical protein - fruit fly (Drosophila melanog
 XM_370992 Homo sapiens LOC388298 (LOC388298), mRNA
 XM_370993 Homo sapiens similar to coenzyme A diphosphatase (LOC388299), mRNA
 XM_370995 Homo sapiens snail homolog 3 (Drosophila) (SNAI3), mRNA
 XM_370997 Homo sapiens similar to Brain-type organic cation transporter (Solute carrier
 XM_371001 Homo sapiens LOC388317 (LOC388317), mRNA
 XM_371006 Homo sapiens similar to RIKEN cDNA C730027E14 (LOC388323), mRNA
 XM_371008 Homo sapiens similar to Death effector filament-forming Ced-4-like apoptos
 XM_371009 Homo sapiens LOC388327 (LOC388327), mRNA
 XM_371010 Homo sapiens hypothetical protein MGC49942 (MGC49942), mRNA
 XM_371012 Homo sapiens similar to ENSANGP00000015193 (LOC388329), mRNA
 XM_371013 Homo sapiens similar to Gag-Pro-Pol protein (LOC388332), mRNA
 XM_371014 Homo sapiens similar to Williams Beuren syndrome chromosome region 19 i
 XM_371015 Homo sapiens ubiquitin specific protease 43 (USP43), mRNA
 XM_371016 Homo sapiens similar to RIKEN cDNA A730055C05 gene (LOC388335), mR
 XM_371017 Homo sapiens similar to hypothetical protein D430041B17 (LOC388336), mF
 XM_371018 Homo sapiens similar to CDRT15 protein (LOC388337), mRNA
 XM_371019 Homo sapiens similar to ribosomal protein (LOC388339), mRNA
 XM_371020 Homo sapiens LOC388341 (LOC388341), mRNA
 XM_371023 Homo sapiens similar to ribosomal protein L13; 60S ribosomal protein L13; b
 XM_371024 Homo sapiens similar to poly(A) binding protein interacting protein 1 isoform
 XM_371026 Homo sapiens similar to hypothetical protein FLJ10847 (LOC388351), mRNA
 XM_371028 Homo sapiens similar to AF038169 protein (LOC388353), mRNA
 XM_371032 Homo sapiens similar to bb329D4.2.1 (novel protein similar to a truncated nu
 XM_371034 Homo sapiens similar to PDZ and LIM domain 1 (elfin); carboxy terminal LIM
 XM_371035 Homo sapiens similar to 40S ribosomal protein S7 (S8) (LOC388363), mRNA
 XM_371036 Homo sapiens KIAA0100 gene product (KIAA0100), mRNA
 XM_371039 Homo sapiens hypothetical protein MGC19764 (MGC19764), mRNA
 XM_371043 Homo sapiens similar to TBC1 domain family member 3 (Rab GTPase-active
 XM_371046 Homo sapiens similar to TBC1 domain family member 3 (Rab GTPase-active
 XM_371052 Homo sapiens Nck, Ash and phospholipase C binding protein (NAP4), mRNA
 XM_371053 Homo sapiens LOC388381 (LOC388381), mRNA
 XM_371054 Homo sapiens LOC388382 (LOC388382), mRNA

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XM_371056 Homo sapiens similar to RIKEN cDNA 4933439F11 (LOC388389), mRNA
 XM_371057 Homo sapiens hypothetical protein LOC201175 (LOC201175), mRNA
 XM_371058 Homo sapiens similar to KIAA0563 gene product (LOC388391), mRNA
 XM_371059 Homo sapiens similar to FALZ protein (LOC388392), mRNA
 XM_371062 Homo sapiens similar to Hypothetical protein KIAA0563 (LOC388395), mRNA
 XM_371063 Homo sapiens similar to KIAA0563 gene product (LOC388396), mRNA
 XM_371066 Homo sapiens similar to SMT3 suppressor of mit two 3 homolog 2 (LOC388397), mRNA
 XM_371067 Homo sapiens similar to Eukaryotic translation initiation factor 4E (eIF4E) (el
 XM_371068 Homo sapiens similar to ribosomal protein L7 (LOC388401), mRNA
 XM_371069 Homo sapiens similar to 40S ribosomal protein S2 (LOC388402), mRNA
 XM_371070 Homo sapiens similar to Yippee-like protein 2 (DisGeorge syndrome-related p
 XM_371074 Homo sapiens putative ankyrin-repeat containing protein (DKFZP564D166),
 XM_371077 Homo sapiens similar to PP905 (LOC388413), mRNA
 XM_371078 Homo sapiens similar to galectin 3 binding protein; L3 antigen; Mac-2-binding
 XM_371079 Homo sapiens Fas binding protein 1 (FBF-1), mRNA
 XM_371081 Homo sapiens LOC388424 (LOC388424), mRNA
 XM_371082 Homo sapiens hypothetical protein FLJ20753 (FLJ20753), mRNA
 XM_371083 Homo sapiens LOC388428 (LOC388428), mRNA
 XM_371084 Homo sapiens KIAA1447 protein (KIAA1447), mRNA
 XM_371085 Homo sapiens hypothetical protein LOC339229 (LOC339229), mRNA
 XM_371086 Homo sapiens similar to dysferlin-interacting protein 1 (LOC388432), mRNA
 XM_371088 Homo sapiens similar to hypothetical protein DKFZp434F142 (LOC388437),
 XM_371089 Homo sapiens similar to MEGF11 protein (LOC388438), mRNA
 XM_371092 Homo sapiens similar to cytokine (LOC388440), mRNA
 XM_371097 Homo sapiens LOC388444 (LOC388444), mRNA
 XM_371098 Homo sapiens hypothetical protein LOC348262 (LOC348262), mRNA
 XM_371105 Homo sapiens LOC388457 (LOC388457), mRNA
 XM_371106 Homo sapiens LOC388458 (LOC388458), mRNA
 XM_371107 Homo sapiens similar to 60S ribosomal protein L6 (TAX-responsive enhance
 XM_371108 Homo sapiens similar to KIAA1314 protein (LOC388462), mRNA
 XM_371109 Homo sapiens hypothetical protein LOC284221 (LOC284221), mRNA
 XM_371110 Homo sapiens similar to acyl-malonyl condensing enzyme (LOC388463), mF
 XM_371111 Homo sapiens similar to breast cancer antigen NY-BR-1.1 (LOC388469), mF
 XM_371113 Homo sapiens similar to 60S RIBOSOMAL PROTEIN L21 (LOC388471), mR
 XM_371114 Homo sapiens formin homology 2 domain containing 3 (FHOD3), mRNA
 XM_371115 Homo sapiens similar to 60S ribosomal protein L7a (Surfeit locus protein 3) (L
 XM_371116 Homo sapiens myosin VB (MYO5B), mRNA
 XM_371117 Homo sapiens similar to Nonhistone chromosomal protein HMG-14 (High-mc
 XM_371118 Homo sapiens similar to serologically defined colon cancer antigen 3 (LOC3
 XM_371120 Homo sapiens thioredoxin-like 4 (TXNL4), mRNA
 XM_371121 Homo sapiens similar to ba476115.3 (novel protein similar to septin) (LOC38
 XM_371122 Homo sapiens similar to Dip1-associated protein C (LOC388488), mRNA
 XM_371125 Homo sapiens similar to KIAA1683 protein (LOC388491), mRNA
 XM_371130 Homo sapiens similar to methyl-CpG binding domain protein 3-like 2 (LOC38
 XM_371132 Homo sapiens FLJ38144 protein (FLJ38144), mRNA
 XM_371134 Homo sapiens similar to complement C3-Q2 (LOC388503), mRNA
 XM_371138 Homo sapiens hypothetical protein LOC284390 (LOC284390), mRNA
 XM_371139 Homo sapiens hypothetical protein FLJ14959 (FLJ14959), mRNA
 XM_371140 Homo sapiens similar to zinc finger protein 433 (LOC388507), mRNA
 XM_371141 Homo sapiens similar to ribosomal protein L17 (LOC388508), mRNA
 XM_371142 Homo sapiens similar to hypothetical protein FLJ38281 (LOC388509), mRN/
 XM_371143 Homo sapiens similar to Asialoglycoprotein receptor 2 (Hepatic lectin H2) (A/
 XM_371145 Homo sapiens similar to Cytochrome P450 4F12 (CYP4F12) (LOC388514),
 XM_371146 Homo sapiens KIAA1683 (KIAA1683), mRNA
 XM_371147 Homo sapiens similar to MOST-1 protein (LOC388517), mRNA
 XM_371150 Homo sapiens zinc finger protein 90 (HTF9) (ZNF90), mRNA
 XM_371151 Homo sapiens similar to 40S ribosomal protein S16 (LOC388519), mRNA

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XM_371152 Homo sapiens zinc finger protein 486 (ZNF486), mRNA
 XM_371153 Homo sapiens similar to hypothetical protein (LOC388521), mRNA
 XM_371154 Homo sapiens similar to Zinc finger protein 429 (LOC388522), mRNA
 XM_371155 Homo sapiens similar to 40S ribosomal protein SA (P40) (34/67 kDa laminin
 XM_371157 Homo sapiens similar to CG14939-PA (LOC388526), mRNA
 XM_371158 Homo sapiens hypothetical protein LOC147991 (LOC147991), mRNA
 XM_371159 Homo sapiens similar to regulator of G protein signaling 9-binding protein; R1
 XM_371160 Homo sapiens similar to 60S ribosomal protein L21 (LOC388532), mRNA
 XM_371161 Homo sapiens similar to KIPV467 (LOC388533), mRNA
 XM_371164 Homo sapiens NYD-SP11 protein (NYD-SP11), mRNA
 XM_371165 Homo sapiens similar to SPRED-3 (LOC388538), mRNA
 XM_371167 Homo sapiens syncollin (SYCN), mRNA
 XM_371170 Homo sapiens similar to Zinc finger protein 216 (LOC388545), mRNA
 XM_371174 Homo sapiens zinc finger protein 283 (ZNF283), mRNA
 XM_371175 Homo sapiens zinc finger protein 229 (ZNF229), mRNA
 XM_371176 Homo sapiens similar to CEACAM5 protein (LOC388550), mRNA
 XM_371177 Homo sapiens similar to cardioembryonic antigen-related cell adhesion mol
 XM_371178 Homo sapiens similar to BC043666 protein (LOC388552), mRNA
 XM_371179 Homo sapiens F-box only protein 34-like (FBXO34L), mRNA
 XM_371181 Homo sapiens nanos homolog 2 (Drosophila) (NANOS2), mRNA
 XM_371182 Homo sapiens similar to BC282485_1 (LOC388554), mRNA
 XM_371183 Homo sapiens similar to RPRC483 (LOC388555), mRNA
 XM_371184 Homo sapiens KIAA1183 protein (KIAA1183), mRNA
 XM_371187 Homo sapiens hypothetical gene MGC45922 (MGC45922), mRNA
 XM_371190 Homo sapiens hypothetical protein LOC162967 (LOC162967), mRNA
 XM_371191 Homo sapiens similar to RIKEN cDNA 1300007C21 (LOC388560), mRNA
 XM_371192 Homo sapiens similar to KIAA2033 protein (LOC388561), mRNA
 XM_371195 Homo sapiens hypothetical protein MGC35045 (MGC35045), mRNA
 XM_371196 Homo sapiens LOC388564 (LOC388564), mRNA
 XM_371197 Homo sapiens similar to zinc finger protein 111 (LOC388565), mRNA
 XM_371198 Homo sapiens similar to Zinc finger protein 471 (EZF1-related protein 1) (LC
 XM_371200 Homo sapiens similar to R30217_1 (LOC388567), mRNA
 XM_371201 Homo sapiens similar to Zinc finger protein 324 (Zinc finger protein ZF5128)
 XM_371202 Homo sapiens CXYorf1-related protein (FLJ00038), mRNA
 XM_371204 Homo sapiens similar to 60S ribosomal protein L23a (LOC388574), mRNA
 XM_371205 Homo sapiens similar to hypothetical protein DKFZp434F142 (LOC388576),
 XM_371206 Homo sapiens similar to F-box only protein 25 (LOC388578), mRNA
 XM_371207 Homo sapiens similar to beta-tubulin 4Q (LOC388579), mRNA
 XM_371208 Homo sapiens similar to RIKEN cDNA 1110035L05 (LOC388581), mRNA
 XM_371210 Homo sapiens taste receptor, type 1, member 3 (TAS1R3), mRNA
 XM_371214 Homo sapiens KIAA0450 gene product (KIAA0450), mRNA
 XM_371215 Homo sapiens similar to hairy and enhancer of split 5 (LOC388585), mRNA
 XM_371216 Homo sapiens LOC388591 (LOC388591), mRNA
 XM_371221 Homo sapiens similar to WD repeat domain 9 isoform A; WD repeat domain
 XM_371222 Homo sapiens similar to Hypothetical protein MGC37938 (LOC388595), mR
 XM_371223 Homo sapiens similar to 40S ribosomal protein S16 (LOC388596), mRNA
 XM_371225 Homo sapiens hypothetical protein LOC284729 (LOC284729), mRNA
 XM_371227 Homo sapiens ciliary rootlet coiled-coil, rootletin (CROCC), mRNA
 XM_371230 Homo sapiens similar to heparan-sulfate 6-sulfotransferase (LOC388605), m
 XM_371232 Homo sapiens similar to Succinate dehydrogenase [ubiquinone] cytochrome
 XM_371234 Homo sapiens similar to RhCE protein (LOC388607), mRNA
 XM_371235 Homo sapiens similar to mutant membrane protein RhCe (LOC388608), mR1
 XM_371236 Homo sapiens hypothetical protein FLJ10747 (FLJ10747), mRNA
 XM_371237 Homo sapiens cation channel, sperm associated 4 (CATSPER4), mRNA
 XM_371238 Homo sapiens LOC388610 (LOC388610), mRNA
 XM_371239 Homo sapiens similar to EAPG6122 (LOC388611), mRNA
 XM_371241 Homo sapiens similar to RIKEN cDNA 9330177P20 (LOC388618), mRNA

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XM_371243 Homo sapiens similar to 60S ribosomal protein L21 (LOC388621), mRNA
 XM_371244 Homo sapiens LOC388624 (LOC388624), mRNA
 XM_371245 Homo sapiens similar to FLJ00406 protein (LOC388625), mRNA
 XM_371246 Homo sapiens hypothetical protein FLJ21156 (FLJ21156), mRNA
 XM_371248 Homo sapiens LOC374973 (LOC374973), mRNA
 XM_371249 Homo sapiens similar to cytochrome P450 4Z1 (LOC388629), mRNA
 XM_371250 Homo sapiens LOC388630 (LOC388630), mRNA
 XM_371252 Homo sapiens similar to heterogeneous nuclear ribonucleoprotein A3 (LOC3)
 XM_371253 Homo sapiens similar to Low-density lipoprotein receptor-related protein 2 pr
 XM_371254 Homo sapiens similar to Ubiquitin carboxyl-terminal hydrolase 24 (Ubiquitin t
 XM_371257 Homo sapiens KIAA1799 protein (KIAA1799), mRNA
 XM_371258 Homo sapiens hypothetical protein FLJ10770 (KIAA1579), mRNA
 XM_371259 Homo sapiens hypothetical protein DKFZp547I048 (DKFZp547I048), mRNA
 XM_371260 Homo sapiens similar to crystallin, zeta; NADPH:quinone reductase (LOC3886)
 XM_371261 Homo sapiens similar to Triosephosphate isomerase (TIM) (LOC388642), m
 XM_371262 Homo sapiens EGF, latrophilin and seven transmembrane domain containing
 XM_371263 Homo sapiens mucolin 2 (MCOLN2), mRNA
 XM_371265 Homo sapiens similar to Guanylate binding protein 4 (LOC388646), mRNA
 XM_371267 Homo sapiens hypothetical protein LOC164045 (LOC164045), mRNA
 XM_371268 Homo sapiens similar to 170002BK03 protein (LOC388649), mRNA
 XM_371269 Homo sapiens similar to RIKEN cDNA 2900024C23 (LOC388650), mRNA
 XM_371273 Homo sapiens similar to 40S ribosomal protein SA (P40) (34/67 kDa laminin
 XM_371276 Homo sapiens similar to hypothetical protein MGC8902 (LOC388658), mRNA
 XM_371277 Homo sapiens similar to RIKEN cDNA C230093N12 (LOC388659), mRNA
 XM_371278 Homo sapiens similar to mitsugumin29 (LOC388660), mRNA
 XM_371279 Homo sapiens amphoterin-induced gene (KIAA1163), mRNA
 XM_371280 Homo sapiens similar to Orphan sodium- and chloride-dependent neurotrans
 XM_371281 Homo sapiens similar to DKFZP564K247 protein (LOC388665), mRNA
 XM_371283 Homo sapiens LOC388666 (LOC388666), mRNA
 XM_371284 Homo sapiens T-box 15 (TBX15), mRNA
 XM_371285 Homo sapiens 3-beta-hydroxysteroid dehydrogenase, tissue-type heart (LOC
 XM_371286 Homo sapiens hypothetical protein MGC45731 (MGC45731), mRNA
 XM_371288 Homo sapiens similar to cyclophilin-LC; cyclophilin homolog overexpressed i
 XM_371291 Homo sapiens similar to hypothetical protein FLJ21308 (LOC388673), mRNA
 XM_371292 Homo sapiens LOC388674 (LOC388674), mRNA
 XM_371299 Homo sapiens similar to KIAA0454 protein (LOC388681), mRNA
 XM_371301 Homo sapiens similar to hypothetical protein FLJ21308 (LOC388685), mRNA
 XM_371302 Homo sapiens similar to cyclophilin-LC; cyclophilin homolog overexpressed i
 XM_371304 Homo sapiens similar to cyclophilin-LC; cyclophilin homolog overexpressed i
 XM_371305 Homo sapiens similar to hypothetical protein KIAA0454 - human (fragment) (L
 XM_371306 Homo sapiens similar to KIAA0454 protein (LOC388689), mRNA
 XM_371310 Homo sapiens similar to hypothetical protein SB145 (LOC388695), mRNA
 XM_371311 Homo sapiens hypothetical protein FLJ36032 (FLJ36032), mRNA
 XM_371312 Homo sapiens hypothetical protein FLJ39117 (FLJ39117), mRNA
 XM_371313 Homo sapiens similar to dJ14N1.2 (novel S-100/CalBP type calcium binding
 XM_371314 Homo sapiens similar to skin-specific protein (LOC388699), mRNA
 XM_371315 Homo sapiens LOC388701 (LOC388701), mRNA
 XM_371316 Homo sapiens similar to 40S ribosomal protein SA (P40) (34/67 kDa laminin
 XM_371320 Homo sapiens FLJ00193 protein (FLJ00193), mRNA
 XM_371326 Homo sapiens similar to Putative dimethylaniline monooxygenase [N-oxide fi
 XM_371328 Homo sapiens hypothetical gene supported by BC007071 (dJ383J4.3), mRNA
 XM_371329 Homo sapiens similar to Protein translation factor SUI1 homolog (Sui1sio1) (L
 XM_371330 Homo sapiens similar to ba92K2.2 (similar to ubiquitin) (LOC388720), mRNA
 XM_371331 Homo sapiens similar to Hypothetical protein CBG13135 (LOC388722), mRNA
 XM_371332 Homo sapiens kinesin family member 21B (KIF21B), mRNA
 XM_371333 Homo sapiens similar to hypothetical protein FLJ37794 (LOC388724), mRNA
 XM_371334 Homo sapiens similar to KIAA1151 protein (LOC388725), mRNA

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XM_371335 Homo sapiens similar to osteotesticular protein tyrosine phosphatase (LOC3
 XM_371336 Homo sapiens similar to alpha tubulin (LOC388728), mRNA
 XM_371338 Homo sapiens hypothetical protein LOC93273 (LOC93273), mRNA
 XM_371339 Homo sapiens similar to 40S ribosomal protein S25 (LOC388733), mRNA
 XM_371340 Homo sapiens similar to Px19-like protein (25 kDa protein of relevant evoluti
 XM_371343 Homo sapiens similar to KIAA0663 protein (LOC388739), mRNA
 XM_371344 Homo sapiens similar to calpain 8 (LOC388743), mRNA
 XM_371346 Homo sapiens hypothetical protein LOC347813 (LOC347813), mRNA
 XM_371350 Homo sapiens similar to RIKEN cDNA 1810063B05 (LOC388753), mRNA
 XM_371351 Homo sapiens similar to 60S ribosomal protein L35 (LOC388754), mRNA
 XM_371352 Homo sapiens formin 2 (FMN2), mRNA
 XM_371353 Homo sapiens LOC388756 (LOC388756), mRNA
 XM_371354 Homo sapiens hypothetical protein FLJ10157 (FLJ10157), mRNA
 XM_371355 Homo sapiens LOC388759 (LOC388759), mRNA
 XM_371356 Homo sapiens similar to olfactory receptor GA_x6K02T2NHDJ-9721756-972
 XM_371357 Homo sapiens similar to Olfactory receptor 2M6 (LOC388762), mRNA
 XM_371358 Homo sapiens olfactory receptor, family 2, subfamily M, member 4 (OR2M4),
 XM_371359 Homo sapiens similar to Ankr3-prov protein (LOC388763), mRNA
 XM_371369 Homo sapiens C219-reactive peptide (FLJ39207), mRNA
 XM_371374 Homo sapiens similar to hypothetical protein (LOC388774), mRNA
 XM_371380 Homo sapiens S100 calcium binding protein A13 (S100A13), mRNA
 XM_371384 Homo sapiens similar to AGO2 (LOC388776), mRNA
 XM_371385 Homo sapiens similar to hypothetical protein DJ528E19.C1.1 (LOC388777),
 XM_371388 Homo sapiens DKFZp434D177-like (DKFZp434D177-like), mRNA
 XM_371390 Homo sapiens similar to hypothetical protein (LOC388783), mRNA
 XM_371391 Homo sapiens similar to dJ680N4.2 (ubiquitin-conjugating enzyme E2D 3 (h
 XM_371395 Homo sapiens similar to dJ1093G12.6 (A novel protein) (LOC388794), mRNA
 XM_371397 Homo sapiens similar to hypothetical protein FLJ33620 (LOC388795), mRN
 XM_371398 Homo sapiens myosin, heavy polypeptide 7B, cardiac muscle, beta (MYH7B)
 XM_371399 Homo sapiens chromosome 20 open reading frame 142 (C20orf142), mRNA
 XM_371401 Homo sapiens chromosome 20 open reading frame 106 (C20orf106), mRNA
 XM_371402 Homo sapiens similar to dJ1153D9.4 (novel protein) (LOC388799), mRNA
 XM_371403 Homo sapiens LOC388802 (LOC388802), mRNA
 XM_371405 Homo sapiens similar to bA476I15.3 (novel protein similar to septin) (LOC38
 XM_371407 Homo sapiens similar to Ankyrin repeat domain protein 18A (LOC388812), n
 XM_371409 Homo sapiens similar to peptidylprolyl isomerase A (cyclophilin A) (LOC388
 XM_371411 Homo sapiens similar to GNGT1 protein (LOC388819), mRNA
 XM_371413 Homo sapiens similar to Protein CGI-27 (C21orf19-like protein) (LOC38882
 XM_371416 Homo sapiens similar to C21orf258 (LOC388828), mRNA
 XM_371417 Homo sapiens KIAA0179 (KIAA0179), mRNA
 XM_371418 Homo sapiens similar to PRED59 (LOC388830), mRNA
 XM_371421 Homo sapiens similar to hypothetical protein (LOC388840), mRNA
 XM_371423 Homo sapiens similar to hypothetical protein DKFZp434P211.1 - human (frag
 XM_371424 Homo sapiens similar to breakpoint cluster region isoform 1 (LOC388847), n
 XM_371429 Homo sapiens similar to hypothetical protein (LOC388852), mRNA
 XM_371430 Homo sapiens similar to sushi domain containing 2; Sushi domain (SCR rep
 XM_371431 Homo sapiens similar to Gamma-glutamyltranspeptidase 1 precursor (Gamr
 XM_371436 Homo sapiens similar to E2F transcription factor 6 isoform a (LOC388861), n
 XM_371437 Homo sapiens similar to dJ831C21.3 (novel protein similar to DKFZP434P21
 XM_371455 Homo sapiens LOC388862 (LOC388862), mRNA
 XM_371459 Homo sapiens hypothetical protein MGC1842 (MGC1842), mRNA
 XM_371460 Homo sapiens LOC388866 (LOC388866), mRNA
 XM_371461 Homo sapiens KIAA1671 protein (KIAA1671), mRNA
 XM_371463 Homo sapiens similar to hypothetical protein 4930562D19 (LOC388891), mF
 XM_371466 Homo sapiens LOC388900 (LOC388900), mRNA
 XM_371468 Homo sapiens hypothetical protein MGC40042 (MGC40042), mRNA
 XM_371469 Homo sapiens LOC388906 (LOC388906), mRNA

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XM_371470 Homo sapiens similar to ribosomal protein L5; 60S ribosomal protein L5 (LOC
 XM_371471 Homo sapiens chromosome 22 open reading frame 1 (C22orf1), mRNA
 XM_371474 Homo sapiens plexin B2 (PLXNB2), mRNA
 XM_371476 Homo sapiens similar to Doublecortin domain-containing protein 2 (RU2S pr
 XM_371477 Homo sapiens similar to RIKEN cDNA 5830483C08 gene (LOC388926), mR
 XM_371478 Homo sapiens LOC388927 (LOC388927), mRNA
 XM_371479 Homo sapiens LOC388928 (LOC388928), mRNA
 XM_371480 Homo sapiens similar to tudor domain containing 6 protein (LOC388929), mF
 XM_371481 Homo sapiens LOC388931 (LOC388931), mRNA
 XM_371484 Homo sapiens selenoprotein 1, 1 (K1AA1724), mRNA
 XM_371485 Homo sapiens similar to RIKEN cDNA 1700001C02 (LOC388936), mRNA
 XM_371486 Homo sapiens similar to phospholipase B (LOC388937), mRNA
 XM_371487 Homo sapiens phospholipase B1 (PLB1), mRNA
 XM_371488 Homo sapiens similar to CDNA sequence BC027072 (LOC388939), mRNA
 XM_371489 Homo sapiens similar to hypothetical protein (L1H 3 region) - human (LOC3E
 XM_371490 Homo sapiens similar to cysteine and histidine-rich domain (CHORD)-contai
 XM_371491 Homo sapiens LOC388946 (LOC388946), mRNA
 XM_371492 Homo sapiens similar to signal-transducing adaptor protein-2; brk kinase sut
 XM_371493 Homo sapiens similar to hypothetical protein (LOC388950), mRNA
 XM_371494 Homo sapiens similar to Testis-specific Y-encoded-like protein 1 (TSPY-like
 XM_371495 Homo sapiens similar to 40S ribosomal protein SA (P40) (34/67 kDa laminin
 XM_371496 Homo sapiens similar to Px19-like protein (25 kDa protein of relevant evoluit
 XM_371497 Homo sapiens similar to expressed sequence C79663 (LOC388957), mRNA
 XM_371500 Homo sapiens similar to hypothetical protein 4921507A12 (LOC388960), mF
 XM_371501 Homo sapiens hypothetical protein MGC22014 (MGC22014), mRNA
 XM_371502 Homo sapiens similar to RIKEN cDNA 1810056O20 (LOC388962), mRNA
 XM_371503 Homo sapiens similar to Retinol dehydrogenase 12 (LOC388963), mRNA
 XM_371504 Homo sapiens similar to hepatitis C virus core-binding protein 6; cervical can
 XM_371505 Homo sapiens similar to Phosphatidylethanolamine-binding protein (PEBP) (L
 XM_371506 Homo sapiens LOC388969 (LOC388969), mRNA
 XM_371511 Homo sapiens similar to anaphase promoting complex subunit 1; anaphase-I
 XM_371513 Homo sapiens similar to RAN-binding protein 2-like 1 isoform 1; sperm mem
 XM_371514 Homo sapiens similar to WW domain binding protein 1 (LOC388975), mRNA
 XM_371515 Homo sapiens similar to protein that is immuno-reactive with anti-PTH polycl
 XM_371517 Homo sapiens similar to immunoglobulin kappa light chain VC region (LOC3I
 XM_371534 Homo sapiens similar to hypothetical protein (LOC389000), mRNA
 XM_371535 Homo sapiens similar to hypothetical protein DKFZp434A171 (LOC389002),
 XM_371536 Homo sapiens similar to tripartite motif-containing 43 (LOC389004), mRNA
 XM_371537 Homo sapiens similar to RING finger protein 18 (Testis-specific ring-finger pr
 XM_371539 Homo sapiens similar to hypothetical protein (LOC389007), mRNA
 XM_371540 Homo sapiens ankyrin-related (UNQ2430), mRNA
 XM_371542 Homo sapiens RW1 protein (RW1), mRNA
 XM_371543 Homo sapiens LOC389012 (LOC389012), mRNA
 XM_371544 Homo sapiens similar to Sodium/hydrogen exchanger 4 (Na⁺)/H⁺ exchang
 XM_371546 Homo sapiens similar to Elongation factor 1-alpha 1 (EF-1-alpha-1) (Elongat
 XM_371547 Homo sapiens similar to Elongation factor 1-alpha 1 (EF-1-alpha-1) (Elongat
 XM_371552 Homo sapiens similar to RAN-binding protein 2-like 1 isoform 1; sperm mem
 XM_371555 Homo sapiens similar to myosin-VIIB (LOC389031), mRNA
 XM_371558 Homo sapiens similar to FKSG30 (LOC389036), mRNA
 XM_371561 Homo sapiens similar to CDNA sequence BC043098 (LOC389039), mRNA
 XM_371564 Homo sapiens similar to sequence-specific single-stranded-DNA-binding pro
 XM_371567 Homo sapiens similar to pote protein; Expressed in prostate, ovary, testis, ar
 XM_371568 Homo sapiens similar to pote protein; Expressed in prostate, ovary, testis, ar
 XM_371569 Homo sapiens similar to Probable mitochondrial import receptor subunit TOB
 XM_371571 Homo sapiens similar to breast cancer antigen NY-BR-1 (LOC389052), mRN
 XM_371572 Homo sapiens similar to heterogenous nuclear ribonucleoprotein K (LOC38
 XM_371573 Homo sapiens neurexophilin 2 (NXP2), mRNA

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XM_371575 Homo sapiens formin binding protein 3 (FNBP3), mRNA
 XM_371576 Homo sapiens KIAA1189 protein (KIAA1189), mRNA
 XM_371577 Homo sapiens similar to RNA polymerase B transcription factor 3 (MGC2390
 XM_371581 Homo sapiens similar to zinc finger protein Sp5 (LOC389058), mRNA
 XM_371583 Homo sapiens LOC389064 (LOC389064), mRNA
 XM_371584 Homo sapiens similar to RIKEN cDNA B830010L13 gene (LOC389065), mR
 XM_371585 Homo sapiens LOC389066 (LOC389066), mRNA
 XM_371586 Homo sapiens hypothetical protein FLJ25415 (FLJ25415), mRNA
 XM_371588 Homo sapiens similar to Selenide,water dikinase 1 (Selenophosphate synthe
 XM_371590 Homo sapiens KIAA1571 protein (KIAA1571), mRNA
 XM_371591 Homo sapiens similar to RIKEN cDNA 9430067K14 gene: Ras GTPase-activ
 XM_371592 Homo sapiens similar to RIKEN cDNA D630023F18 (LOC389073), mRNA
 XM_371593 Homo sapiens similar to REGULATED ENDOCRINE SPECIFIC PROTEIN 11
 XM_371594 Homo sapiens similar to CAVP-target protein (CAVPT) (LOC389076), mRNA
 XM_371595 Homo sapiens dedicator of cytokinesis 10 (DOCK10), mRNA
 XM_371600 Homo sapiens similar to enterokinase (LOC389083), mRNA
 XM_371603 Homo sapiens similar to hypothetical protein FLJ40243 (LOC389085), mRN/
 XM_371604 Homo sapiens hypothetical protein FLJ37034 (FLJ37034), mRNA
 XM_371605 Homo sapiens hypothetical protein LOC151174 (LOC151174), mRNA
 XM_371606 Homo sapiens similar to seven transmembrane helix receptor (LOC389090),
 XM_371614 Homo sapiens hypothetical protein FLJ10707 (FLJ10707), mRNA
 XM_371617 Homo sapiens TBP-interacting protein (TIP120B), mRNA
 XM_371618 Homo sapiens similar to nucleoporin 210; nuclear pore membrane glycoprot
 XM_371619 Homo sapiens FYVE, RhoGEF and PH domain containing 5 (FGD5), mRNA
 XM_371621 Homo sapiens similar to RIKEN cDNA B830010L13 gene (LOC389099), mR/
 XM_371622 Homo sapiens similar to 60S ribosomal protein L23a (LOC389101), mRNA
 XM_371623 Homo sapiens similar to YPLR6490 (LOC389102), mRNA
 XM_371625 Homo sapiens LOC389106 (LOC389106), mRNA
 XM_371626 Homo sapiens similar to Hsp70/Hsp90 organizing protein homolog CG2720-I
 XM_371629 Homo sapiens similar to RIKEN cDNA 1110038F21 (LOC389111), mRNA
 XM_371630 Homo sapiens similar to ribosomal protein S27 (LOC389112), mRNA
 XM_371631 Homo sapiens similar to hypothetical protein FLJ35863 (LOC389114), mRN/
 XM_371632 Homo sapiens FLJ36157 protein (FLJ36157), mRNA
 XM_371638 Homo sapiens similar to mouse fat 1 cadherin (LOC389117), mRNA
 XM_371639 Homo sapiens similar to VLLR9392 (LOC389118), mRNA
 XM_371641 Homo sapiens similar to RIKEN cDNA 4921517D21 (LOC389120), mRNA
 XM_371643 Homo sapiens similar to hypothetical protein MGC39725 (LOC389124), mR/
 XM_371645 Homo sapiens similar to 40S ribosomal protein S10 (LOC389127), mRNA
 XM_371647 Homo sapiens similar to CG9996-PA (LOC389129), mRNA
 XM_371649 Homo sapiens similar to hypothetical protein SB153 isoform 1 (LOC389130),
 XM_371653 Homo sapiens similar to hypothetical protein FLJ11292 (LOC389135), mRN/
 XM_371655 Homo sapiens similar to SVH-B (LOC389137), mRNA
 XM_371656 Homo sapiens similar to olfactory receptor GA_x54KRFPKG5P-55348161-55
 XM_371658 Homo sapiens similar to laminin receptor 1 (ribosomal protein SA); P40-3, fur
 XM_371660 Homo sapiens similar to seven transmembrane helix receptor (LOC389144),
 XM_371662 Homo sapiens hypothetical protein LOC255330 (LOC255330), mRNA
 XM_371663 Homo sapiens similar to WD repeat domain 10 isoform 3 (LOC389147), mR/
 XM_371664 Homo sapiens KIAA1257 protein (KIAA1257), mRNA
 XM_371665 Homo sapiens LOC389151 (LOC389151), mRNA
 XM_371666 Homo sapiens LOC389152 (LOC389152), mRNA
 XM_371668 Homo sapiens similar to 60S ribosomal protein L21 (LOC389156), mRNA
 XM_371670 Homo sapiens similar to Plscr1 protein (LOC389158), mRNA
 XM_371671 Homo sapiens similar to Chromosome 1 open reading frame 37 (LOC389164
 XM_371672 Homo sapiens similar to template acylating factor-I alpha (LOC389168), mR/
 XM_371674 Homo sapiens similar to hypothetical protein FLJ14957 (LOC389170), mRN/
 XM_371677 Homo sapiens similar to phosphoserine aminotransferase isoform 1 (LOC38/
 XM_371678 Homo sapiens LOC389174 (LOC389174), mRNA

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XM_371679 Homo sapiens similar to ribosomal protein L22 (LOC389175), mRNA
 XM_371680 Homo sapiens LOC389177 (LOC389177), mRNA
 XM_371681 Homo sapiens similar to RING finger protein 13 (LOC389178), mRNA
 XM_371682 Homo sapiens similar to 5-hydroxytryptamine receptor 3 subunit C (LOC389179), mRNA
 XM_371683 Homo sapiens LOC389187 (LOC389187), mRNA
 XM_371684 Homo sapiens similar to FSHD Region Gene 2 protein (LOC389192), mRNA
 XM_371687 Homo sapiens LOC389197 (LOC389197), mRNA
 XM_371690 Homo sapiens LOC389202 (LOC389202), mRNA
 XM_371691 Homo sapiens LOC389203 (LOC389203), mRNA
 XM_371692 Homo sapiens similar to LINE-1 REVERSE TRANSCRIPTASE HOMOLOG 1 (LOC389204), mRNA
 XM_371693 Homo sapiens LOC389206 (LOC389206), mRNA
 XM_371694 Homo sapiens similar to ENSANGP0000012385 (LOC389207), mRNA
 XM_371695 Homo sapiens similar to RIKEN cDNA 4732406D01 gene (LOC389208), mRNA
 XM_371697 Homo sapiens similar to expressed sequence AW060714 (LOC389211), mRNA
 XM_371698 Homo sapiens similar to KIAA1680 protein (LOC401145), mRNA
 XM_371701 Homo sapiens similar to template acylating factor-1 alpha (LOC389217), mRNA
 XM_371702 Homo sapiens similar to Ribosomal protein L7A CG3314-PD (LOC389218), mRNA
 XM_371705 Homo sapiens LOC389221 (LOC389221), mRNA
 XM_371706 Homo sapiens hypothetical protein KIAA1109 (KIAA1109), mRNA
 XM_371709 Homo sapiens ring finger protein 150 (RNF150), mRNA
 XM_371710 Homo sapiens LOC389227 (LOC389227), mRNA
 XM_371711 Homo sapiens similar to GRIK2 protein (LOC389228), mRNA
 XM_371714 Homo sapiens similar to ring finger protein 129 (LOC389239), mRNA
 XM_371715 Homo sapiens similar to alpha NAC1.9.2, protein (LOC389240), mRNA
 XM_371717 Homo sapiens odd Oz/Ten-m homolog 3 (ODZ3), mRNA
 XM_371718 Homo sapiens similar to vesicle-associated soluble NSF attachment protein 1 (LOC389241), mRNA
 XM_371719 Homo sapiens similar to RST1689 (LOC389255), mRNA
 XM_371722 Homo sapiens similar to Leudine-rich repeat-containing protein 14 (LOC389256), mRNA
 XM_371725 Homo sapiens similar to hypothetical protein (LOC389261), mRNA
 XM_371726 Homo sapiens similar to antifreeze glycoprotein precursor - black rockcod (LOC389262), mRNA
 XM_371728 Homo sapiens similar to CDNA sequence BC012016 (LOC389276), mRNA
 XM_371729 Homo sapiens similar to RPE-spondin (LOC389279), mRNA
 XM_371731 Homo sapiens similar to bA110H4.2 (similar to membrane protein) (LOC389280), mRNA
 XM_371732 Homo sapiens similar to RIKEN cDNA C230086A09 gene (LOC389282), mRNA
 XM_371733 Homo sapiens hypothetical protein FLJ23577 (FLJ23577), mRNA
 XM_371736 Homo sapiens similar to FKSG62 (LOC389286), mRNA
 XM_371738 Homo sapiens similar to annexin II receptor (LOC389289), mRNA
 XM_371740 Homo sapiens hypothetical protein FLJ23563 (FLJ23563), mRNA
 XM_371741 Homo sapiens similar to hypothetical protein (LOC389293), mRNA
 XM_371743 Homo sapiens similar to bA110H4.2 (similar to membrane protein) (LOC389294), mRNA
 XM_371749 Homo sapiens similar to bA110H4.2 (similar to membrane protein) (LOC389295), mRNA
 XM_371754 Homo sapiens similar to nonhistone chromosomal protein HMG-17 - rat (LOC389296), mRNA
 XM_371755 Homo sapiens similar to Rho-guanine nucleotide exchange factor (Rho-Inter) (LOC389297), mRNA
 XM_371757 Homo sapiens similar to 60S ribosomal protein L7 (LOC389305), mRNA
 XM_371758 Homo sapiens similar to ribosomal protein L10a (LOC389308), mRNA
 XM_371759 Homo sapiens hypothetical protein FLJ11292 (FLJ11292), mRNA
 XM_371760 Homo sapiens hypothetical protein LOC116068 (LOC116068), mRNA
 XM_371761 Homo sapiens KIAA0825 protein (KIAA0825), mRNA
 XM_371762 Homo sapiens similar to Proteasome activator complex subunit 2 (Proteasom) (LOC389310), mRNA
 XM_371763 Homo sapiens LOC389313 (LOC389313), mRNA
 XM_371764 Homo sapiens similar to NADH dehydrogenase subunit 5 (LOC389314), mRNA
 XM_371765 Homo sapiens LOC389315 (LOC389315), mRNA
 XM_371768 Homo sapiens LOC389320 (LOC389320), mRNA
 XM_371769 Homo sapiens LOC389321 (LOC389321), mRNA
 XM_371770 Homo sapiens similar to heterogeneous nuclear ribonucleoprotein K (LOC389322), mRNA
 XM_371771 Homo sapiens similar to hypothetical protein (LOC389323), mRNA
 XM_371772 Homo sapiens LOC389326 (LOC389326), mRNA

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XM_371776 Homo sapiens similar to hypothetical protein 4933429F08 (LOC389337), mRNA
 XM_371777 Homo sapiens hypothetical protein LOC348938 (LOC348938), mRNA
 XM_371778 Homo sapiens similar to RIKEN cDNA 4921536K21 (LOC389341), mRNA
 XM_371781 Homo sapiens similar to 60S ribosomal protein L10 (QM protein homolog) (L
 XM_371782 Homo sapiens similar to hypothetical protein (LOC389346), mRNA
 XM_371783 Homo sapiens similar to NY-REN-7 antigen (LOC389347), mRNA
 XM_371790 Homo sapiens similar to 6-pyruvoyl-tetrahydropterin synthase (LOC389351),
 XM_371791 Homo sapiens similar to High mobility group protein 4 (HMG-4) (High mobility
 XM_371793 Homo sapiens similar to bA47615.3 (novel protein similar to septin) (LOC38
 XM_371796 Homo sapiens similar to Protein phosphatase 1, regulatory subunit 3D (Prot
 XM_371797 Homo sapiens LOC389365 (LOC389365), mRNA
 XM_371798 Homo sapiens similar to C6orf52 protein (LOC389366), mRNA
 XM_371801 Homo sapiens O-acyltransferase (membrane bound) domain containing 1 (O
 XM_371809 Homo sapiens chromosome 6 open reading frame 205 (C6orf205), mRNA
 XM_371812 Homo sapiens major histocompatibility complex, class II, DQ alpha 1 (HLA-D
 XM_371813 Homo sapiens kinesin family member C1 (KIFC1), mRNA
 XM_371814 Homo sapiens similar to Rps15a protein (LOC389382), mRNA
 XM_371815 Homo sapiens similar to AAL3045 (LOC389383), mRNA
 XM_371816 Homo sapiens similar to RIKEN cDNA 4930539E08 (LOC389384), mRNA
 XM_371817 Homo sapiens similar to Cytosol aminopeptidase (Leucine aminopeptidase) i
 XM_371818 Homo sapiens similar to Cytosol aminopeptidase (Leucine aminopeptidase) i
 XM_371819 Homo sapiens similar to 60S ribosomal protein L12 (LOC389387), mRNA
 XM_371820 Homo sapiens similar to hypothetical protein BC006605 (LOC389389), mRNA
 XM_371822 Homo sapiens chromosome 6 open reading frame 110 (C6orf110), mRNA
 XM_371823 Homo sapiens similar to hypothetical protein DKFZp434D2328 (LOC389394)
 XM_371824 Homo sapiens similar to heterogeneous nuclear ribonucleoprotein A3 (LOC3
 XM_371825 Homo sapiens similar to BXMAS2-10 protein (LOC389396), mRNA
 XM_371826 Homo sapiens similar to IVF19356 (LOC389400), mRNA
 XM_371829 Homo sapiens similar to ENSANGP00000009924 (LOC389405), mRNA
 XM_371832 Homo sapiens KIAA1411 (KIAA1411), mRNA
 XM_371835 Homo sapiens inhibitor of Bruton agammaglobulinemia tyrosine kinase (IBTK
 XM_371837 Homo sapiens similar to oxidoreductase UCPA (LOC389416), mRNA
 XM_371838 Homo sapiens ubiquitin specific protease 45 (USP45), mRNA
 XM_371841 Homo sapiens similar to hypothetical protein (L1H 3 region) - human (LOC3E
 XM_371842 Homo sapiens LOC389421 (LOC389421), mRNA
 XM_371843 Homo sapiens similar to ribosomal protein S27a (LOC389425), mRNA
 XM_371844 Homo sapiens TSPY-like (TSPYL), mRNA
 XM_371845 Homo sapiens similar to MGC32805 protein (LOC389427), mRNA
 XM_371846 Homo sapiens similar to ribosomal protein L5; 60S ribosomal protein L5 (Lo
 XM_371847 Homo sapiens similar to RIKEN cDNA 2310057J18 (LOC389429), mRNA
 XM_371848 Homo sapiens chromosome 6 open reading frame 115 (C6orf115), mRNA
 XM_371849 Homo sapiens chromosome 6 open reading frame 198 (C6orf198), mRNA
 XM_371850 Homo sapiens similar to hypothetical protein 9130014G24 (LOC389431), mF
 XM_371851 Homo sapiens similar to RIKEN cDNA E130306M17 gene (LOC389432), mF
 XM_371853 Homo sapiens similar to 60S ribosomal protein L27a (LOC389435), mRNA
 XM_371856 Homo sapiens similar to frazzled C6G581-PA (LOC389444), mRNA
 XM_371857 Homo sapiens similar to 60S ribosomal protein L21 (LOC389445), mRNA
 XM_371858 Homo sapiens similar to apolipoprotein(a) (EC 3.4.21.-) - rhesus macaque (fi
 XM_371863 Homo sapiens family with sequence similarity 20, member C (FAM20C), mR
 XM_371873 Homo sapiens similar to zinc finger protein 316; kruppel-related zinc finger p
 XM_371874 Homo sapiens similar to Matn2-prov protein (LOC389462), mRNA
 XM_371877 Homo sapiens KIAA0960 protein (KIAA0960), mRNA
 XM_371878 Homo sapiens hypothetical protein FLJ14712 (FLJ14712), mRNA
 XM_371879 Homo sapiens LOC389466 (LOC389466), mRNA
 XM_371884 Homo sapiens similar to 40S ribosomal protein S26 (LOC389472), mRNA
 XM_371885 Homo sapiens similar to Neuronal protein 3.1 (p311 protein) (LOC389473), n
 XM_371889 Homo sapiens similar to RP9 protein (LOC389477), mRNA

XM_371891 Homo sapiens KIAA0877 protein (KIAA0877), mRNA
 XM_371897 Homo sapiens hypothetical protein DKFZp76112123 (DKFZp76112123), mRNA
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 XM_371904 Homo sapiens similar to Protein p8 (Candidate of metastasis 1) (LOC389492)
 XM_371917 Homo sapiens similar to Williams Beuren syndrome chromosome region 19 (LOC371923)
 XM_371923 Homo sapiens similar to opposite strand transcription unit to Stag3; Gats pro
 XM_371925 Homo sapiens similar to transcription factor GTF2IRD2 (LOC389524), mRNA
 XM_371930 Homo sapiens similar to PMS4 homolog mismatch repair protein - human (LOC371933)
 XM_371933 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 9A (PPP1R19A)
 XM_371936 Homo sapiens similar to hypothetical protein MGC56855 (LOC389538), mRNA
 XM_371939 Homo sapiens similar to CG14977-PA (LOC389541), mRNA
 XM_371943 Homo sapiens similar to Williams-Beuren syndrome critical region protein 19 (LOC371948)
 XM_371948 Homo sapiens similar to zinc finger protein 312; forebrain embryonic zinc finger protein 312 (LOC389550), mRNA
 XM_371949 Homo sapiens similar to cardiac telomodlin (LOC389550), mRNA
 XM_371953 Homo sapiens KIAA1466 protein (KIAA1466), mRNA
 XM_371954 Homo sapiens nucleoporin 205kDa (NUP205), mRNA
 XM_371956 Homo sapiens KIAA1549 protein (KIAA1549), mRNA
 XM_371960 Homo sapiens KIAA1277 (KIAA1277), mRNA
 XM_371995 Homo sapiens similar to hypothetical protein MGC41943 (LOC389592), mRNA
 XM_372002 Homo sapiens similar to amyotrophic lateral sclerosis 2 (juvenile) chromosome 2 (LOC372004)
 XM_372004 Homo sapiens LOC389602 (LOC389602), mRNA
 XM_372005 Homo sapiens LOC389603 (LOC389603), mRNA
 XM_372006 Homo sapiens similar to vasoactive Intestinal peptide receptor 2 (LOC389606)
 XM_372009 Homo sapiens similar to bA476115.3 (novel protein similar to septin) (LOC389607)
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 XM_372011 Homo sapiens similar to HARL2754 (LOC389610), mRNA
 XM_372013 Homo sapiens similar to FLJ10408 protein (LOC389611), mRNA
 XM_372017 Homo sapiens similar to seven transmembrane helix receptor (LOC389616), mRNA
 XM_372018 Homo sapiens similar to hypothetical protein FLJ10408 (LOC389617), mRNA
 XM_372019 Homo sapiens similar to FLJ10408 protein (LOC389618), mRNA
 XM_372024 Homo sapiens LOC389622 (LOC389622), mRNA
 XM_372025 Homo sapiens similar to Activated RNA polymerase II transcriptional coactivator 1 (LOC389626)
 XM_372026 Homo sapiens similar to seven transmembrane helix receptor (LOC389626), mRNA
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 XM_372030 Homo sapiens similar to FLJ10408 protein (LOC389633), mRNA
 XM_372031 Homo sapiens mitochondrial tumor suppressor gene 1 (MTSG1), mRNA
 XM_372035 Homo sapiens LOC389643 (LOC389643), mRNA
 XM_372036 Homo sapiens similar to 60S ribosomal protein L5 (LOC389644), mRNA
 XM_372037 Homo sapiens LOC389649 (LOC389649), mRNA
 XM_372038 Homo sapiens hypothetical protein FLJ32731 (FLJ32731), mRNA
 XM_372039 Homo sapiens similar to hypothetical protein (L1H 3 region) - human (LOC389650)
 XM_372040 Homo sapiens similar to asparagine synthetase; glutamine-dependent asparagine synthetase (LOC389658), mRNA
 XM_372041 Homo sapiens similar to RPLK9433 (LOC389658), mRNA
 XM_372042 Homo sapiens similar to polycystin 1-like 3 (LOC389660), mRNA
 XM_372045 Homo sapiens similar to hypothetical protein (LOC389663), mRNA
 XM_372046 Homo sapiens similar to tropomyosin 4 (LOC389667), mRNA
 XM_372047 Homo sapiens similar to hypothetical protein FLJ10307 (LOC389668), mRNA
 XM_372048 Homo sapiens similar to 40S ribosomal protein SA (P40) (34/67 kDa laminin)
 XM_372050 Homo sapiens similar to Heterogeneous nuclear ribonucleoprotein A1 (Helix-Loop-Helix) (LOC389678)
 XM_372054 Homo sapiens LOC389678 (LOC389678), mRNA
 XM_372055 Homo sapiens similar to hypothetical protein (LOC389679), mRNA
 XM_372058 Homo sapiens Pvt1 oncogene homolog, MYC activator (mouse) (PVT1), mRNA
 XM_372060 Homo sapiens similar to FLJ46354 protein (LOC389690), mRNA
 XM_372062 Homo sapiens similar to FLJ46354 protein (LOC389694), mRNA
 XM_372063 Homo sapiens similar to epiplakin (LOC389697), mRNA
 XM_372069 Homo sapiens similar to prot GOR (LOC389699), mRNA

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XM_372074 Homo sapiens similar to Selenoprotein T (LOC389704), mRNA
 XM_372075 Homo sapiens similar to RIKEN cDNA 3110001D03 gene (M. musculus) (MC
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 XM_372077 Homo sapiens LOC389708 (LOC389708), mRNA
 XM_372078 Homo sapiens similar to hypothetical protein (L1H 3 region) - human (LOC38
 XM_372083 Homo sapiens KIAA1161 (KIAA1161), mRNA
 XM_372086 Homo sapiens similar to RAB1B, member RAS oncogene family; small GTP-
 XM_372088 Homo sapiens similar to cell recognition molecule CASPR3 isoform 1 (LOC3
 XM_372089 Homo sapiens similar to chromosome 9 open reading frame 36 (LOC389723
 XM_372090 Homo sapiens similar to cell recognition molecule CASPR3 isoform 1 (LOC3
 XM_372092 Homo sapiens similar to FK506-binding protein 4 (Peptidyl-prolyl cis-trans is
 XM_372094 Homo sapiens similar to chromosome 9 open reading frame 36 (LOC389730
 XM_372097 Homo sapiens similar to cell recognition molecule CASPR3 isoform 1 (LOC3
 XM_372099 Homo sapiens similar to bA251017.4 (similar to methylentetrahydrofolate d
 XM_372100 Homo sapiens LOC389739 (LOC389739), mRNA
 XM_372102 Homo sapiens LOC389742 (LOC389742), mRNA
 XM_372103 Homo sapiens LOC389744 (LOC389744), mRNA
 XM_372104 Homo sapiens similar to breast cancer antigen NY-BR-1 (LOC389745), mRN
 XM_372108 Homo sapiens similar to Ribosome biogenesis protein BMS1 homolog (LOC
 XM_372109 Homo sapiens similar to mitochondrial C1-tetrahydrofolate synthase (LOC38
 XM_372110 Homo sapiens aquaporin 7-like (LOC375719), mRNA
 XM_372111 Homo sapiens similar to mitochondrial C1-tetrahydrofolate synthase (LOC38
 XM_372112 Homo sapiens similar to phosphoglucosyltransferase 5 (LOC389753), mRNA
 XM_372114 Homo sapiens similar to mitochondrial C1-tetrahydrofolate synthase (LOC38
 XM_372116 Homo sapiens similar to COBW domain containing protein 3 (LOC389760), r
 XM_372117 Homo sapiens similar to CG3073-PA (LOC375743), mRNA
 XM_372118 Homo sapiens similar to RIKEN cDNA 1700013B16 (LOC389761), mRNA
 XM_372119 Homo sapiens LOC389762 (LOC389762), mRNA
 XM_372120 Homo sapiens similar to RIKEN cDNA 1700013B16 (LOC389763), mRNA
 XM_372121 Homo sapiens similar to kinesin family member 27 (LOC389764), mRNA
 XM_372122 Homo sapiens similar to kinesin family member 27 (LOC389765), mRNA
 XM_372123 Homo sapiens similar to RIKEN cDNA 4921517D22 (LOC389766), mRNA
 XM_372124 Homo sapiens zinc finger, CCHC domain containing 6 (ZCCHC6), mRNA
 XM_372125 Homo sapiens similar to potassium channel tetramerisation domain contain
 XM_372128 Homo sapiens similar to Osteotesticular phosphatase; protein tyrosine phosph
 XM_372132 Homo sapiens similar to RIKEN cDNA 2810453I06 (LOC389776), mRNA
 XM_372133 Homo sapiens KIAA1529 (KIAA1529), mRNA
 XM_372137 Homo sapiens similar to RIKEN cDNA 4732481H14 (LOC389785), mRNA
 XM_372138 Homo sapiens LOC389786 (LOC389786), mRNA
 XM_372140 Homo sapiens LOC389789 (LOC389789), mRNA
 XM_372141 Homo sapiens similar to phosphatidylinositol phosphate kinase-like protein (I
 XM_372142 Homo sapiens LOC389791 (LOC389791), mRNA
 XM_372143 Homo sapiens hypothetical protein LOC375757 (LOC375757), mRNA
 XM_372148 Homo sapiens similar to CDNA sequence BC034076 (LOC389796), mRNA
 XM_372150 Homo sapiens LOC389799 (LOC389799), mRNA
 XM_372154 Homo sapiens similar to bA74P14.2 (novel protein) (LOC389803), mRNA
 XM_372157 Homo sapiens similar to HSPC324 (LOC389811), mRNA
 XM_372159 Homo sapiens similar to CG15216-PA (LOC389813), mRNA
 XM_372160 Homo sapiens similar to LPAL6438 (LOC389814), mRNA
 XM_372161 Homo sapiens similar to CDNA sequence BC004853 (LOC389816), mRNA
 XM_372162 Homo sapiens LOC389817 (LOC389817), mRNA
 XM_372163 Homo sapiens similar to cell recognition molecule CASPR3 isoform 1 (LOC3
 XM_372168 Homo sapiens LOC389821 (LOC389821), mRNA
 XM_372169 Homo sapiens similar to hypothetical protein (LOC389822), mRNA
 XM_372175 Homo sapiens LOC389823 (LOC389823), mRNA
 XM_372177 Homo sapiens similar to Surfteil locus protein 1 (LOC389825), mRNA
 XM_372180 Homo sapiens similar to RIKEN cDNA 1110002H13 (LOC389827), mRNA

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XM_372181 Homo sapiens similar to MSTP058 (LOC389828), mRNA
 XM_372182 Homo sapiens hypothetical protein FLJ39378 (FLJ39378), mRNA
 XM_372186 Homo sapiens similar to hypothetical protein MGC27019 (LOC389830), mRNA
 XM_372190 Homo sapiens similar to AF038169 protein (LOC389832), mRNA
 XM_372191 Homo sapiens similar to hypothetical protein MGC27019 (LOC389833), mRNA
 XM_372192 Homo sapiens similar to RIKEN cDNA 2700049P18 (LOC389835), mRNA
 XM_372193 Homo sapiens KIAA1751 protein (KIAA1751), mRNA
 XM_372194 Homo sapiens hypothetical protein MGC13275 (MGC13275), mRNA
 XM_372195 Homo sapiens agrin (AGRN), mRNA
 XM_372197 Homo sapiens similar to Glutathione peroxidase 1 (GSPX-1) (Cellular glutat
 XM_372198 Homo sapiens hypothetical protein MGC17403 (MGC17403), mRNA
 XM_372199 Homo sapiens similar to MAP/ERK kinase kinase 5; apoptosis signal regulat
 XM_372200 Homo sapiens similar to Ran-specific GTPase-activating protein (Ran bindin
 XM_372201 Homo sapiens similar to hypothetical protein FLJ35782 (LOC389843), mRNA
 XM_372202 Homo sapiens similar to ferritin, heavy polypeptide-like 17 (LOC389844), mR
 XM_372203 Homo sapiens similar to Amine oxidase [flavin-containing] A (Monoamine oxi
 XM_372204 Homo sapiens similar to 40S ribosomal protein SA (P40) (34/67 kDa laminin
 XM_372205 Homo sapiens similar to hypothetical protein, MGC:7199 (LOC389850), mRNA
 XM_372208 Homo sapiens similar to dJ54B20.3 (novel protein similar to lysoczyme C (1,4
 XM_372209 Homo sapiens similar to Acetyl-coenzyme A acyltransferase 2 (LOC389854),
 XM_372210 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 3F (PPP
 XM_372212 Homo sapiens similar to G antigen 7 (LOC389855), mRNA
 XM_372213 Homo sapiens similar to ubiquitin specific protease 27, X chromosome (LOC
 XM_372214 Homo sapiens similar to nuclear protein p30 (LOC389857), mRNA
 XM_372223 Homo sapiens similar to 28S ribosomal protein S18c, mitochondrial precursor
 XM_372224 Homo sapiens similar to PAGE-5 protein (LOC389860), mRNA
 XM_372226 Homo sapiens similar to Zinc finger X-linked protein ZXDB (LOC389862), mR
 XM_372227 Homo sapiens similar to PAL-1 mRNA-binding protein; chromodomain helic
 XM_372231 Homo sapiens hypothetical protein FLJ20105 (FLJ20105), mRNA
 XM_372233 Homo sapiens similar to Selenide,water dikinase 1 (Selenophosphate synthe
 XM_372239 Homo sapiens hypothetical protein BC007652 (LOC92129), mRNA
 XM_372245 Homo sapiens similar to R28830_1 (LOC389885), mRNA
 XM_372247 Homo sapiens similar to RIKEN cDNA 1110012O05 (LOC389887), mRNA
 XM_372248 Homo sapiens similar to FLJ20527 protein (LOC389888), mRNA
 XM_372253 Homo sapiens similar to ENSANGP00000013187 (LOC389891), mRNA
 XM_372254 Homo sapiens similar to Mothers against decapentaplegic homolog interacti
 XM_372255 Homo sapiens similar to MGC38553 protein (LOC389895), mRNA
 XM_372257 Homo sapiens similar to ubiquitin-conjugating enzyme E2N (LOC389898), m
 XM_372258 Homo sapiens similar to HS1 binding protein (LOC389899), mRNA
 XM_372261 Homo sapiens similar to DKFZP586L0724 protein (LOC389900), mRNA
 XM_372262 Homo sapiens similar to ATP-dependent DNA helicase II, 70 kDa subunit (L
 XM_372267 Homo sapiens zinc finger protein 275 (ZNF275), mRNA
 XM_372268 Homo sapiens similar to Extracellular matrix protein 2 precursor (Matrix glyco
 XM_372272 Homo sapiens LOC389905 (LOC389905), mRNA
 XM_372273 Homo sapiens similar to Serine/threonine protein kinase PRKX (Protein kinas
 XM_372274 Homo sapiens similar to Serine/threonine protein kinase PRKX (Protein kinas
 XM_372275 Homo sapiens LOC389911 (LOC389911), mRNA
 XM_372282 Homo sapiens cytokine receptor-like factor 2 (CRLF2), mRNA
 XM_372286 Homo sapiens neuroligin 4, Y linked (NLGN4Y), mRNA
 XM_372289 Homo sapiens chromosome Y open reading frame 15A (Ycorf15A), mRNA
 XM_372292 Homo sapiens similar to RNA binding motif protein, Y chromosome, family 1
 XM_372295 Homo sapiens similar to bA476115.3 (novel protein similar to septin) (LOC38
 XM_372296 Homo sapiens similar to deleted in azoospermia (LOC389926), mRNA
 XM_372302 Homo sapiens similar to 3-hydroxyhexobarbital dehydrogenase 1/3-alpha, 17
 XM_372303 Homo sapiens similar to proline-rich proteoglycan 2 (LOC389936), mRNA
 XM_372304 Homo sapiens similar to St:dz211013.2 (novel protein) (LOC389938), mRNA
 XM_372305 Homo sapiens similar to Gliacolin (LOC389941), mRNA

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XM_372443 Homo sapiens similar to Olfactory receptor 8A1 (OST025) (LOC390275), mRNA
 XM_372444 Homo sapiens similar to Bcl-2 homologous antagonist/killer (Apoptosis regul
 XM_372445 Homo sapiens similar to retinitis pigmentosa GTPase regulator (LOC390278
 XM_372447 Homo sapiens similar to eukaryotic translation initiation factor 3, subunit 5 (e
 XM_372448 Homo sapiens similar to signal recognition particle 14kDa (homologous Alu F
 XM_372449 Homo sapiens similar to olfactory receptor GA_x6K02T2PVTD-14054886-14
 XM_372450 Homo sapiens similar to Heterogeneous nuclear ribonucleoprotein A1 (Helix-
 XM_372452 Homo sapiens similar to Peptidylprolyl isomerase A (cyclophilin A) (LOC3902
 XM_372453 Homo sapiens similar to RIKEN cDNA 110014F12 (LOC390300), mRNA
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 XM_372457 Homo sapiens similar to olfactory receptor like protein (LOC390313), mRNA
 XM_372459 Homo sapiens similar to olfactory receptor MOR111-1 (LOC390321), mRNA
 XM_372460 Homo sapiens similar to olfactory receptor MOR112-1 (LOC390323), mRNA
 XM_372461 Homo sapiens similar to olfactory receptor MOR109-1 (LOC390324), mRNA
 XM_372462 Homo sapiens similar to olfactory receptor GA_x6K02T2PULF-11304679-11;
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 XM_372465 Homo sapiens similar to olfactory receptor MOR114-12 (LOC390328), mRNA
 XM_372466 Homo sapiens similar to poly(A) binding protein, cytoplasmic 4 isoform 2 (LO
 XM_372468 Homo sapiens similar to T-box transcription factor TBX20 (LOC390338), mRNA
 XM_372469 Homo sapiens similar to Gag-Pro-Pol-Env protein (LOC390342), mRNA
 XM_372470 Homo sapiens similar to microtubule-associated proteins 1A/1B light chain 3
 XM_372471 Homo sapiens similar to 60S ribosomal protein L10 (QM protein) (Tumor sup
 XM_372472 Homo sapiens similar to Small nuclear ribonucleoprotein Sm D2 (snRNP con
 XM_372473 Homo sapiens similar to eukaryotic translation elongation factor 1 alpha 1; C
 XM_372474 Homo sapiens similar to ribosomal protein L9 (LOC390353), mRNA
 XM_372476 Homo sapiens similar to dynein, axonemal, heavy chain 11; situs inversus vi
 XM_372480 Homo sapiens similar to KIAA1786 protein (LOC390365), mRNA
 XM_372482 Homo sapiens similar to 40S ribosomal protein S6 (Phosphoprotein NP33) (L
 XM_372483 Homo sapiens similar to proline rich protein 2 (LOC390371), mRNA
 XM_372485 Homo sapiens similar to Mitochondrial ornithine transporter 1 (Solute carrier
 XM_372486 Homo sapiens similar to Alpha-N-acetyl-neuraminy-2,3-beta-galactosyl-1,3-
 XM_372487 Homo sapiens similar to chromosome 9 open reading frame 12; 1,3,4,5,6-pe
 XM_372488 Homo sapiens similar to histone 1, H2bc; H2B histone family, member S (LO
 XM_372490 Homo sapiens similar to Striatin (LOC390387), mRNA
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 XM_372493 Homo sapiens similar to ADP/ATP carrier protein T2 - human (LOC390405),
 XM_372494 Homo sapiens similar to chromosome 15 open reading frame 2 (LOC390414
 XM_372496 Homo sapiens similar to ADP-ribosylation factor 4 (LOC390423), mRNA
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 XM_372498 Homo sapiens LOC390427 (LOC390427), mRNA
 XM_372499 Homo sapiens similar to Olfactory receptor 4M1 (LOC390428), mRNA
 XM_372500 Homo sapiens similar to Olfactory receptor 4N2 (LOC390429), mRNA
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 XM_372506 Homo sapiens similar to Olfactory receptor 11G2 (LOC390439), mRNA
 XM_372507 Homo sapiens similar to Olfactory receptor 11H4 (LOC390442), mRNA
 XM_372508 Homo sapiens similar to Ribonuclease-like protein 9 precursor (LOC390443)
 XM_372509 Homo sapiens similar to Olfactory receptor 5AU1 (LOC390445), mRNA
 XM_372521 Homo sapiens similar to SMT3 suppressor of mif two 3 homolog 2 (LOC3904
 XM_372522 Homo sapiens similar to Plasmodium falciparum trophozoite antigen r45-like
 XM_372524 Homo sapiens LOC390481 (LOC390481), mRNA
 XM_372525 Homo sapiens similar to ribosomal protein L31 (LOC390485), mRNA
 XM_372527 Homo sapiens similar to 60S ribosomal protein L21 (LOC390488), mRNA

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XM_372528 Homo sapiens hypothetical protein FLJ36749 (FLJ36749), mRNA
 XM_372532 Homo sapiens similar to Alpha-1-antitrypsin-related protein precursor (LOC3:
 XM_372534 Homo sapiens similar to proline-rich glycoprotein (sgp158) (LOC390507), mR
 XM_372535 Homo sapiens similar to MGC53446 protein (LOC390508), mRNA
 XM_372536 Homo sapiens similar to chloride intracellular channel 6; chloride channel fo
 XM_372542 Homo sapiens similar to protein kinase related to Raf protein kinases; Metho
 XM_372543 Homo sapiens similar to immunoglobulin heavy-chain-2 light-chain-2 VH seg
 XM_372544 Homo sapiens similar to hect domain and RLD 2 (LOC390533), mRNA
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 XM_372573 Homo sapiens similar to hypothetical protein BC009980 (LOC390594), mRN
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 XM_372575 Homo sapiens similar to cell division cycle 2-like 1 (PITSLRE proteins) isofor
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 XM_372598 Homo sapiens similar to Olfactory receptor 4F14 (LOC390650), mRNA
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 XM_372617 Homo sapiens similar to Protein C20orf27 (LOC390690), mRNA
 XM_372618 Homo sapiens similar to Filamin B (FLN-B) (Beta-filamin) (Actin-binding like

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XM_372622 Homo sapiens similar to hypothetical protein MGC5244 (LOC390697), mRNA
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 XM_372629 Homo sapiens similar to Ig H-chain V-region (DP-40) (LOC390710), mRNA
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 XM_372689 Homo sapiens similar to mitochondrial carrier triple repeat 1 (LOC390828), m
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 XM_372704 Homo sapiens similar to ribosomal protein L35 (LOC390876), mRNA
 XM_372705 Homo sapiens similar to adenylate kinase (EC 2.7.4.3), cytosolic - common c

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XM_372785 Homo sapiens similar to peptidylprolyl isomerase A (cyclophilin A) (LOC391C
 XM_372788 Homo sapiens similar to SNAG1 (LOC391086), mRNA
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 XM_372882 Homo sapiens hypothetical protein LOC128954 (LOC128954), mRNA

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XM_372884 Homo sapiens cat eye syndrome chromosome region, candidate 2 (CECR2),
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 XM_372928 Homo sapiens similar to C-terminal binding protein 2 isoform 2; ribeye (LOC:
 XM_372933 Homo sapiens similar to Immunoglobulin kappa light chain variable region O
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 XM_372964 Homo sapiens similar to ATP synthase alpha chain, mitochondrial precursor
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 XM_372973 Homo sapiens similar to guanidinoacetate methyltransferase; GAMT (LOC39
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XM_372981 Homo sapiens similar to Hypothetical protein CBG20540 (LOC391501), mRNA
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XM_373084 Homo sapiens similar to ribosomal protein L31 (LOC391789), mRNA
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 XM_373087 Homo sapiens similar to cadherin 12, type 2 preproprotein; Br-cadherin; cadl
 XM_373088 Homo sapiens similar to cadherin 12, type 2 preproprotein; Br-cadherin; cadl
 XM_373089 Homo sapiens similar to hypothetical protein FLJ10891 (LOC391796), mRNA
 XM_373090 Homo sapiens LOC391797 (LOC391797), mRNA
 XM_373091 Homo sapiens similar to 60S ribosomal protein L10 (QM protein homolog) (L
 XM_373092 Homo sapiens similar to peptidyl-Pro cis trans isomerase (LOC391807), mR
 XM_373093 Homo sapiens similar to 28 kDa heat- and acid-stable phosphoprotein (PDG
 XM_373094 Homo sapiens similar to protease (prosome, macropain) 26S subunit, ATPase
 XM_373096 Homo sapiens similar to 60S RIBOSOMAL PROTEIN L21 (LOC391823), mR
 XM_373097 Homo sapiens similar to Thyroid hormone receptor-associated protein compl
 XM_373098 Homo sapiens similar to beta-tropomyosin (LOC391844), mRNA
 XM_373099 Homo sapiens similar to hypothetical protein (LOC391847), mRNA
 XM_373100 Homo sapiens similar to 60S ribosomal protein L10 (QM protein homolog) (L
 XM_373101 Homo sapiens similar to neuralized homolog (LOC391849), mRNA
 XM_373103 Homo sapiens similar to Beta-1,4-galactosyltransferase 7 (Beta-1,4-GalTase
 XM_373106 Homo sapiens similar to Tb-291 membrane associated protein (LOC391859)
 XM_373107 Homo sapiens similar to cAMP-regulated phosphoprotein 19 (ARPP-19) (LO
 XM_373109 Homo sapiens hypothetical protein LOC340156 (LOC340156), mRNA
 XM_373156 Homo sapiens hypothetical protein MGC26484 (MGC26484), mRNA
 XM_373157 Homo sapiens similar to Phosphorylase B kinase gamma catalytic chain, ske
 XM_373170 Homo sapiens hypothetical protein LOC54103 (LOC54103), mRNA
 XM_373171 Homo sapiens similar to dynein, cytoplasmic, light peptide; 8kD LC; dynein L
 XM_373178 Homo sapiens similar to Zn-alpha-2-glycoprotein precursor (LOC392082), m
 XM_373180 Homo sapiens similar to Cohesin subunit SA-3 (Stromal antigen 3) (SCC3 hc
 XM_373190 Homo sapiens similar to dJ753D5.2 (novel protein similar to RPS17 (40S rib
 XM_373209 Homo sapiens similar to seven transmembrane helix receptor (LOC392133),
 XM_373210 Homo sapiens similar to olfactory receptor MOR261-1 (LOC392138), mRNA
 XM_373214 Homo sapiens similar to Mtr3 (mRNA transport regulator 3)-homolog; Mtr3 (n
 XM_373219 Homo sapiens similar to gastrulation brain homeobox 1 (LOC392152), mRNA/
 XM_373223 Homo sapiens similar to polycystic kidney disease 1-like 3 (LOC392159), mF
 XM_373224 Homo sapiens similar to cAMP-dependent protein kinase type I-beta regulat
 XM_373227 Homo sapiens similar to Olfactory receptor 4F3 (LOC392167), mRNA
 XM_373233 Homo sapiens similar to 60S ribosomal protein L10 (QM protein) (Tumor sup
 XM_373234 Homo sapiens similar to demidefensin 3 (LOC392182), mRNA
 XM_373236 Homo sapiens similar to chromosome 11 open reading frame2; chromosome
 XM_373238 Homo sapiens similar to deubiquitinating enzyme 3 (LOC392188), mRNA
 XM_373239 Homo sapiens similar to ubiquitin-specific protease 17-like protein (LOC3921
 XM_373241 Homo sapiens similar to beta-1,4-mannosyltransferase; beta-1,4 mannosyltr
 XM_373242 Homo sapiens similar to 60S ribosomal protein L10 (QM protein homolog) (L
 XM_373243 Homo sapiens similar to deubiquitinating enzyme 3 (LOC392197), mRNA
 XM_373245 Homo sapiens similar to 60S ribosomal protein L32 (LOC392202), mRNA
 XM_373246 Homo sapiens similar to Ac2-210 (LOC392208), mRNA
 XM_373248 Homo sapiens similar to VENT-like homeobox 2; hemopoietic progenitor hon
 XM_373249 Homo sapiens similar to Ig lambda light chain leader and V-region (LOC3922
 XM_373250 Homo sapiens similar to Band 4.1-like protein 5 (LOC392218), mRNA
 XM_373252 Homo sapiens similar to coiled-coil-helix-coiled-coil-helix domain containi
 XM_373253 Homo sapiens similar to RIKEN cDNA 0610038D11 (LOC392222), mRNA
 XM_373255 Homo sapiens similar to hypothetical class II basic helix-loop-helix protein (L
 XM_373257 Homo sapiens similar to large subunit ribosomal protein L36a (LOC392248),
 XM_373259 Homo sapiens similar to glyceraldehyde-3-phosphate dehydrogenase (LOC3
 XM_373260 Homo sapiens similar to growth differentiation factor 16 (LOC392255), mRNA/
 XM_373263 Homo sapiens similar to laminin-binding protein (LOC392262), mRNA
 XM_373265 Homo sapiens similar to cDNA sequence BC024139 (LOC392274), mRNA
 XM_373266 Homo sapiens similar to sphingomyelin phosphodiesterase 3, neutral membr
 XM_373270 Homo sapiens similar to Rab coupling protein; Rab-interacting recycling prot

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XM_373275 Homo sapiens similar to peptidyl-Pro cis trans isomerase (LOC392285), mRI
 XM_373276 Homo sapiens similar to chromosome 15 open reading frame 12; mitochondri
 XM_373277 Homo sapiens similar to microtubule-associated proteins 1A/1B light chain 3
 XM_373281 Homo sapiens similar to Chloride intracellular channel protein 4 (Intracellular
 XM_373283 Homo sapiens similar to Ferritin light chain (Ferritin L subunit) (LOC392299),
 XM_373284 Homo sapiens similar to High mobility group protein 4-like (HMG-4L) (LOC39
 XM_373287 Homo sapiens similar to RIKEN cDNA 4930412F15 gene (LOC392307), mRI
 XM_373288 Homo sapiens similar to Olfactory receptor 70 (LOC392308), mRNA
 XM_373289 Homo sapiens similar to Olfactory receptor 13J1 (LOC392309), mRNA
 XM_373290 Homo sapiens similar to high mobility group protein homolog HMG4 (LOC39
 XM_373295 Homo sapiens similar to hypothetical protein MGC17986 (LOC392332), mRI
 XM_373297 Homo sapiens LOC392339 (LOC392339), mRNA
 XM_373298 Homo sapiens similar to adenylate kinase 3 alpha like (LOC392347), mRNA
 XM_373300 Homo sapiens similar to TMD1 (LOC392351), mRNA
 XM_373301 Homo sapiens similar to peptidyl-Pro cis trans isomerase (LOC392352), mRI
 XM_373302 Homo sapiens similar to ORF2 (LOC392355), mRNA
 XM_373303 Homo sapiens similar to Eukaryotic translation initiation factor 3 subunit 1 (el
 XM_373304 Homo sapiens similar to Cathepsin L precursor (Major excreted protein) (MEI
 XM_373305 Homo sapiens similar to chromosome 15 open reading frame 2 (LOC392364
 XM_373308 Homo sapiens similar to Olfactory receptor 13C2 (LOC392376), mRNA
 XM_373309 Homo sapiens similar to constitutive photomorphogenic protein (LOC392379
 XM_373310 Homo sapiens similar to large subunit ribosomal protein L36a (LOC392381),
 XM_373311 Homo sapiens similar to ribosomal protein L31 (LOC392382), mRNA
 XM_373312 Homo sapiens similar to 60S ribosomal protein L32 (LOC392384), mRNA
 XM_373313 Homo sapiens olfactory receptor, family 1, subfamily J, member 5 (OR1J5), r
 XM_373314 Homo sapiens similar to G protein-coupled receptor homolog clone H8 (LOC
 XM_373315 Homo sapiens similar to Olfactory receptor 1 L6 (LOC392390), mRNA
 XM_373316 Homo sapiens similar to Olfactory receptor 5 C1 (Olfactory receptor 9-F) (OR
 XM_373317 Homo sapiens similar to Olfactory receptor 1 K1 (LOC392392), mRNA
 XM_373319 Homo sapiens similar to hemicentin (LOC392395), mRNA
 XM_373320 Homo sapiens similar to Von Ebners gland protein precursor (VEG protein) (r
 XM_373322 Homo sapiens similar to Putative MUP-like ilpocalin precursor (LOC392399),
 XM_373324 Homo sapiens similar to Ankyrin repeat domain protein 5 (LOC392404), mRI
 XM_373325 Homo sapiens similar to Pim1 (LOC392405), mRNA
 XM_373326 Homo sapiens similar to hypothetical protein (LOC392406), mRNA
 XM_373327 Homo sapiens similar to Glutamate [NMDA] receptor subunit zeta 1 precursor
 XM_373328 Homo sapiens similar to hypothetical protein FLJ20433 (LOC392409), mRN
 XM_373329 Homo sapiens similar to NADPH-dependent FMN and FAD containing oxidor
 XM_373331 Homo sapiens similar to paranemln (LOC392411), mRNA
 XM_373334 Homo sapiens similar to MGC43306 protein (LOC392413), mRNA
 XM_373335 Homo sapiens similar to Phenylalanine-4-hydroxylase (PAH) (Phe-4-monoox
 XM_373336 Homo sapiens similar to 1060P11.3 (killer inhibitory receptor 2-2-2 (KIR222)
 XM_373337 Homo sapiens similar to protein kinase C, zeta (LOC392420), mRNA
 XM_373338 Homo sapiens similar to bA92K2.2 (similar to ubiquitin) (LOC392425), mRN
 XM_373339 Homo sapiens similar to Nucleolar phosphoprotein p130 (Nucleolar 130 kDa
 XM_373340 Homo sapiens similar to melanoma antigen, family B, 6 (LOC392433), mRN
 XM_373341 Homo sapiens similar to PR264/SC35 (LOC392439), mRNA
 XM_373343 Homo sapiens similar to 60S ribosomal protein L32 (LOC392447), mRNA
 XM_373344 Homo sapiens similar to CpG binding protein (Protein containing PHD finger
 XM_373345 Homo sapiens similar to synovial sarcoma, X breakpoint 6 (LOC392462), mF
 XM_373347 Homo sapiens similar to RIKEN cDNA 201001H14 (LOC392465), mRNA
 XM_373348 Homo sapiens similar to melanoma antigen (LOC392468), mRNA
 XM_373349 Homo sapiens similar to melanoma antigen, family B, 4; melanoma-associat
 XM_373350 Homo sapiens similar to nuclear protein p30 (LOC392468), mRNA
 XM_373351 Homo sapiens similar to dJ83A16.1 (similar to PGAM) (LOC392473), mRN
 XM_373352 Homo sapiens similar to XAGE-5 protein (LOC392475), mRNA
 XM_373353 Homo sapiens similar to Probable G protein-coupled receptor GPR83 precu

XM_373354 Homo sapiens similar to ribosomal protein L31 (LOC392487), mRNA
 XM_373355 Homo sapiens similar to Ras homolog gene family, member G (LOC392489)
 XM_373356 Homo sapiens similar to Translationally controlled tumor protein (TCTP) (p23
 XM_373357 Homo sapiens LOC392491 (LOC392491), mRNA
 XM_373358 Homo sapiens similar to hypothetical protein DKFpZp761H079 isoform 1 (LOC
 XM_373359 Homo sapiens similar to histone H2B-related protein (LOC392512), mRNA
 XM_373362 Homo sapiens similar to dJ820B18.1 (similar to nuclear cap binding protein)
 XM_373365 Homo sapiens similar to Serine/threonine-protein kinase PRP4 homolog (PR
 XM_373366 Homo sapiens similar to beta-tubulin 1 (LOC392528), mRNA
 XM_373367 Homo sapiens similar to RIKEN cDNA 2900070E19 (LOC392531), mRNA
 XM_373368 Homo sapiens similar to Histone H3.3 (LOC392533), mRNA
 XM_373369 Homo sapiens similar to Heat shock cognate 71 kDa protein (LOC392535), r
 XM_373370 Homo sapiens similar to TJP4 protein (LOC392539), mRNA
 XM_373372 Homo sapiens similar to hypothetical protein FLJ20527 (LOC392546), mRNA
 XM_373373 Homo sapiens similar to Glyceraldehyde 3-phosphate dehydrogenase, liver (L
 XM_373374 Homo sapiens similar to Ras-related protein Rab-28 (Rab-26) (LOC392551),
 XM_373378 Homo sapiens similar to hypothetical protein MGC15827 (LOC392559), mR
 XM_373381 Homo sapiens H2A histone family, member B (H2AFB), mRNA
 XM_373382 Homo sapiens similar to neurofilament-like protein (LOC392563), mRNA
 XM_373384 Homo sapiens similar to testis expressed sequence 13A (LOC392566), mRN
 XM_373388 Homo sapiens similar to Menkes Disease (ATP7A) (LOC392570), mRNA
 XM_373391 Homo sapiens similar to Zinc finger protein ZFD25 (LOC392576), mRNA
 XM_373395 Homo sapiens similar to testis specific protein, Y-linked (LOC392582), mRN/
 XM_373397 Homo sapiens similar to testis specific protein, Y-linked (LOC392584), mRN/
 XM_373398 Homo sapiens similar to Testis-Specific Protein Y (TSPY) (LOC392586), mR
 XM_373399 Homo sapiens similar to Sedlin (Trafficking protein particle complex protein 2
 XM_373407 Homo sapiens nuclear receptor subfamily 2, group F, member 6 (NR2F6), m
 XM_373413 Homo sapiens hypothetical protein FLJ20847 (FLJ20847), mRNA
 XM_373419 Homo sapiens valyl-tRNA synthetase 2-like (VARSL), mRNA
 XM_373431 Homo sapiens hypothetical protein BC004360 (LOC87769), mRNA
 XM_373433 Homo sapiens hypothetical protein BC002926 (LOC90379), mRNA
 XM_373440 Homo sapiens chondroitin sulfate synthase 3 (CSS3), mRNA
 XM_373444 Homo sapiens LOC399709 (LOC387630), mRNA
 XM_373445 Homo sapiens LOC399710 (LOC387631), mRNA
 XM_373446 Homo sapiens LOC399711 (LOC387632), mRNA
 XM_373447 Homo sapiens LOC399714 (LOC387633), mRNA
 XM_373448 Homo sapiens LOC399722 (LOC387636), mRNA
 XM_373449 Homo sapiens LOC399724 (LOC387638), mRNA
 XM_373450 Homo sapiens LOC399725 (LOC387639), mRNA
 XM_373451 Homo sapiens hypothetical gene supported by AK056080; AL832706; BC01-
 XM_373452 Homo sapiens LOC399735 (LOC387649), mRNA
 XM_373453 Homo sapiens LOC399741 (LOC387654), mRNA
 XM_373454 Homo sapiens LOC399743 (LOC387656), mRNA
 XM_373456 Homo sapiens hypothetical gene supported by BX538120 (LOC387662), mR
 XM_373460 Homo sapiens LOC399764 (LOC387675), mRNA
 XM_373461 Homo sapiens LOC399775 (LOC387683), mRNA
 XM_373463 Homo sapiens LOC399779 (LOC387686), mRNA
 XM_373464 Homo sapiens LOC399781 (LOC387687), mRNA
 XM_373465 Homo sapiens LOC399784 (LOC387689), mRNA
 XM_373466 Homo sapiens LOC399787 (LOC387690), mRNA
 XM_373468 Homo sapiens LOC399794 (LOC387697), mRNA
 XM_373469 Homo sapiens LOC399797 (LOC387701), mRNA
 XM_373470 Homo sapiens LOC399801 (LOC387705), mRNA
 XM_373471 Homo sapiens LOC399802 (LOC387706), mRNA
 XM_373472 Homo sapiens LOC399805 (LOC387709), mRNA
 XM_373474 Homo sapiens LOC399808 (LOC387710), mRNA
 XM_373475 Homo sapiens LOC399809 (LOC387711), mRNA

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XM_373477 Homo sapiens LOC399813 (LOC387715), mRNA
 XM_373478 Homo sapiens LOC399817 (LOC387717), mRNA
 XM_373479 Homo sapiens LOC399825 (LOC387722), mRNA
 XM_373480 Homo sapiens LOC399828 (LOC387724), mRNA
 XM_373481 Homo sapiens hypothetical gene supported by BC027847; BC047942 (LOC387725), mRNA
 XM_373483 Homo sapiens LOC399847 (LOC387736), mRNA
 XM_373484 Homo sapiens LOC399848 (LOC387737), mRNA
 XM_373485 Homo sapiens LOC399849 (LOC387739), mRNA
 XM_373486 Homo sapiens hypothetical gene supported by AK124823 (LOC387741), mRNA
 XM_373487 Homo sapiens hypothetical gene supported by AK124823 (LOC387742), mRNA
 XM_373488 Homo sapiens LOC399854 (LOC387743), mRNA
 XM_373489 Homo sapiens LOC399856 (LOC387746), mRNA
 XM_373490 Homo sapiens LOC399868 (LOC387754), mRNA
 XM_373491 Homo sapiens LOC399871 (LOC387756), mRNA
 XM_373492 Homo sapiens LOC399874 (LOC387757), mRNA
 XM_373494 Homo sapiens LOC399878 (LOC387760), mRNA
 XM_373495 Homo sapiens LOC399880 (LOC387761), mRNA
 XM_373496 Homo sapiens LOC399882 (LOC387762), mRNA
 XM_373497 Homo sapiens hypothetical gene supported by BC052560 (LOC387763), mRNA
 XM_373498 Homo sapiens LOC399885 (LOC387764), mRNA
 XM_373499 Homo sapiens LOC399887 (LOC387765), mRNA
 XM_373500 Homo sapiens LOC399896 (LOC387771), mRNA
 XM_373502 Homo sapiens LOC399902 (LOC387776), mRNA
 XM_373503 Homo sapiens LOC399903 (LOC387777), mRNA
 XM_373505 Homo sapiens LOC399910 (LOC387783), mRNA
 XM_373506 Homo sapiens LOC399911 (LOC387784), mRNA
 XM_373507 Homo sapiens LOC399916 (LOC387785), mRNA
 XM_373509 Homo sapiens hypothetical gene supported by BC065746 (LOC387789), mRNA
 XM_373511 Homo sapiens LOC399934 (LOC387796), mRNA
 XM_373512 Homo sapiens LOC399935 (LOC387797), mRNA
 XM_373513 Homo sapiens LOC399952 (LOC387810), mRNA
 XM_373515 Homo sapiens LOC399958 (LOC387814), mRNA
 XM_373516 Homo sapiens LOC399966 (LOC387817), mRNA
 XM_373517 Homo sapiens LOC399970 (LOC387819), mRNA
 XM_373518 Homo sapiens hypothetical gene supported by BC039676 (LOC387821), mRNA
 XM_373519 Homo sapiens LOC399976 (LOC387823), mRNA
 XM_373520 Homo sapiens LOC399977 (LOC387824), mRNA
 XM_373521 Homo sapiens LOC399989 (LOC387826), mRNA
 XM_373522 Homo sapiens hypothetical protein LOC283314 (LOC283314), mRNA
 XM_373524 Homo sapiens LOC399997 (LOC387833), mRNA
 XM_373525 Homo sapiens LOC400005 (LOC387838), mRNA
 XM_373527 Homo sapiens LOC400016 (LOC387846), mRNA
 XM_373529 Homo sapiens LOC400017 (LOC387848), mRNA
 XM_373530 Homo sapiens LOC400028 (LOC387853), mRNA
 XM_373531 Homo sapiens LOC400029 (LOC387854), mRNA
 XM_373533 Homo sapiens LOC400034 (LOC387859), mRNA
 XM_373534 Homo sapiens LOC400042 (LOC387863), mRNA
 XM_373535 Homo sapiens LOC400044 (LOC387864), mRNA
 XM_373537 Homo sapiens LOC400049 (LOC387866), mRNA
 XM_373538 Homo sapiens LOC400054 (LOC387869), mRNA
 XM_373539 Homo sapiens LOC400057 (LOC387871), mRNA
 XM_373540 Homo sapiens hypothetical gene supported by BC044741 (LOC387872), mRNA
 XM_373541 Homo sapiens LOC400060 (LOC387873), mRNA
 XM_373542 Homo sapiens LOC400062 (LOC387875), mRNA
 XM_373543 Homo sapiens LOC400063 (LOC387876), mRNA
 XM_373544 Homo sapiens hypothetical gene supported by AK056999; BC013920 (LOC387877), mRNA
 XM_373545 Homo sapiens LOC400071 (LOC387883), mRNA

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XM_373546 Homo sapiens LOC400072 (LOC387884), mRNA
 XM_373547 Homo sapiens LOC400074 (LOC387885), mRNA
 XM_373548 Homo sapiens LOC400075 (LOC387886), mRNA
 XM_373549 Homo sapiens LOC400076 (LOC387887), mRNA
 XM_373550 Homo sapiens LOC400078 (LOC387888), mRNA
 XM_373553 Homo sapiens hypothetical gene supported by BC040060 (LOC387895), mR
 XM_373554 Homo sapiens LOC400089 (LOC387896), mRNA
 XM_373555 Homo sapiens LOC400090 (LOC387897), mRNA
 XM_373556 Homo sapiens LOC400091 (LOC387898), mRNA
 XM_373557 Homo sapiens LOC400100 (LOC387901), mRNA
 XM_373558 Homo sapiens LOC400102 (LOC387903), mRNA
 XM_373559 Homo sapiens LOC400104 (LOC387905), mRNA
 XM_373560 Homo sapiens LOC400107 (LOC387910), mRNA
 XM_373561 Homo sapiens hypothetical gene supported by AK000246 (LOC387915), mR
 XM_373562 Homo sapiens LOC400116 (LOC387917), mRNA
 XM_373563 Homo sapiens LOC400117 (LOC387918), mRNA
 XM_373564 Homo sapiens LOC400133 (LOC387925), mRNA
 XM_373566 Homo sapiens LOC400138 (LOC387931), mRNA
 XM_373567 Homo sapiens LOC400139 (LOC387932), mRNA
 XM_373568 Homo sapiens LOC400143 (LOC387935), mRNA
 XM_373569 Homo sapiens LOC400147 (LOC387937), mRNA
 XM_373570 Homo sapiens LOC400149 (LOC387939), mRNA
 XM_373571 Homo sapiens LOC400150 (LOC387940), mRNA
 XM_373572 Homo sapiens LOC400152 (LOC387941), mRNA
 XM_373573 Homo sapiens LOC400153 (LOC387942), mRNA
 XM_373574 Homo sapiens LOC400155 (LOC387943), mRNA
 XM_373575 Homo sapiens LOC400159 (LOC387946), mRNA
 XM_373576 Homo sapiens LOC400160 (LOC387947), mRNA
 XM_373577 Homo sapiens LOC400162 (LOC387948), mRNA
 XM_373578 Homo sapiens LOC400168 (LOC387950), mRNA
 XM_373580 Homo sapiens hypothetical protein LOC254028 (LOC254028), mRNA
 XM_373581 Homo sapiens LOC400176 (LOC387957), mRNA
 XM_373583 Homo sapiens LOC400196 (LOC387974), mRNA
 XM_373585 Homo sapiens LOC400198 (LOC387977), mRNA
 XM_373587 Homo sapiens LOC400202 (LOC387981), mRNA
 XM_373588 Homo sapiens LOC400204 (LOC387983), mRNA
 XM_373593 Homo sapiens LOC400215 (LOC387989), mRNA
 XM_373594 Homo sapiens LOC400220 (LOC387992), mRNA
 XM_373595 Homo sapiens hypothetical gene supported by BC033546 (LOC387993), mR
 XM_373596 Homo sapiens LOC400225 (LOC387994), mRNA
 XM_373599 Homo sapiens LOC400232 (LOC388001), mRNA
 XM_373600 Homo sapiens LOC400233 (LOC388002), mRNA
 XM_373601 Homo sapiens LOC400235 (LOC388003), mRNA
 XM_373602 Homo sapiens LOC400240 (LOC388006), mRNA
 XM_373603 Homo sapiens chromosome 14 open reading frame 86 (C14orf86), mRNA
 XM_373604 Homo sapiens hypothetical gene supported by BC019017 (LOC388009), mR
 XM_373605 Homo sapiens LOC400245 (LOC388010), mRNA
 XM_373606 Homo sapiens hypothetical gene supported by AK091668 (LOC388011), mR
 XM_373607 Homo sapiens LOC400248 (LOC388014), mRNA
 XM_373608 Homo sapiens LOC400251 (LOC388016), mRNA
 XM_373609 Homo sapiens LOC400252 (LOC388017), mRNA
 XM_373610 Homo sapiens LOC400253 (LOC388018), mRNA
 XM_373611 Homo sapiens LOC400254 (LOC388019), mRNA
 XM_373615 Homo sapiens similar to HSPC053 (LOC388066), mRNA
 XM_373616 Homo sapiens hypothetical gene supported by AK056084; NM_152518 (LOC
 XM_373617 Homo sapiens LOC400321 (LOC388081), mRNA
 XM_373622 Homo sapiens LOC400330 (LOC388091), mRNA

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XM_373623 Homo sapiens LOC400337 (LOC388096), mRNA
 XM_373624 Homo sapiens hypothetical protein LOC283697 (LOC283697), mRNA
 XM_373625 Homo sapiens hypothetical gene supported by BC035481 (LOC388113), mF
 XM_373626 Homo sapiens LOC400382 (LOC388114), mRNA
 XM_373627 Homo sapiens LOC400365 (LOC388117), mRNA
 XM_373628 Homo sapiens LOC400375 (LOC388123), mRNA
 XM_373630 Homo sapiens LOC400379 (LOC388126), mRNA
 XM_373631 Homo sapiens hypothetical gene supported by BC033162 (LOC388129), mR
 XM_373632 Homo sapiens LOC400384 (LOC388131), mRNA
 XM_373633 Homo sapiens LOC400390 (LOC388134), mRNA
 XM_373634 Homo sapiens LOC400394 (LOC388136), mRNA
 XM_373635 Homo sapiens LOC400401 (LOC388139), mRNA
 XM_373636 Homo sapiens LOC400402 (LOC388140), mRNA
 XM_373637 Homo sapiens LOC400404 (LOC388141), mRNA
 XM_373638 Homo sapiens LOC400407 (LOC388142), mRNA
 XM_373639 Homo sapiens LOC400412 (LOC388144), mRNA
 XM_373641 Homo sapiens LOC400425 (LOC388158), mRNA
 XM_373645 Homo sapiens LOC400440 (LOC388169), mRNA
 XM_373646 Homo sapiens LOC400441 (LOC388170), mRNA
 XM_373647 Homo sapiens LOC400443 (LOC388171), mRNA
 XM_373648 Homo sapiens LOC400449 (LOC388176), mRNA
 XM_373649 Homo sapiens LOC400457 (LOC388179), mRNA
 XM_373650 Homo sapiens LOC400458 (LOC388180), mRNA
 XM_373653 Homo sapiens hypothetical gene supported by BC054509 (LOC388193), mF
 XM_373654 Homo sapiens hypothetical gene supported by BC028568 (LOC388197), mR
 XM_373655 Homo sapiens LOC400485 (LOC388198), mRNA
 XM_373657 Homo sapiens LOC400489 (LOC388201), mRNA
 XM_373659 Homo sapiens LOC400491 (LOC388204), mRNA
 XM_373660 Homo sapiens LOC400494 (LOC388206), mRNA
 XM_373662 Homo sapiens LOC400495 (LOC388208), mRNA
 XM_373666 Homo sapiens LOC400501 (LOC388214), mRNA
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XM_374178 Homo sapiens LOC389420 (LOC389420), mRNA
XM_374179 Homo sapiens LOC389422 (LOC389422), mRNA
XM_374180 Homo sapiens LOC389423 (LOC389423), mRNA
XM_374181 Homo sapiens LOC389426 (LOC389426), mRNA
XM_374183 Homo sapiens LOC389433 (LOC389433), mRNA
XM_374184 Homo sapiens LOC389436 (LOC389436), mRNA
XM_374185 Homo sapiens LOC389437 (LOC389437), mRNA
XM_374186 Homo sapiens LOC389438 (LOC389438), mRNA
XM_374187 Homo sapiens LOC389439 (LOC389439), mRNA
XM_374188 Homo sapiens LOC389440 (LOC389440), mRNA
XM_374189 Homo sapiens LOC389441 (LOC389441), mRNA
XM_374190 Homo sapiens LOC389442 (LOC389442), mRNA

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XM_374191 Homo sapiens LOC389443 (LOC389443), mRNA
 XM_374192 Homo sapiens LOC389447 (LOC389447), mRNA
 XM_374193 Homo sapiens LOC389449 (LOC389449), mRNA
 XM_374194 Homo sapiens LOC389450 (LOC389450), mRNA
 XM_374196 Homo sapiens LOC389452 (LOC389452), mRNA
 XM_374198 Homo sapiens LOC389455 (LOC389455), mRNA
 XM_374199 Homo sapiens LOC389456 (LOC389456), mRNA
 XM_374200 Homo sapiens LOC389457 (LOC389457), mRNA
 XM_374202 Homo sapiens similar to Zinc finger protein 268 (Zinc finger protein HZF3) (L
 XM_374203 Homo sapiens LOC389469 (LOC389469), mRNA
 XM_374204 Homo sapiens LOC389470 (LOC389470), mRNA
 XM_374218 Homo sapiens LOC389506 (LOC389506), mRNA
 XM_374222 Homo sapiens LOC389526 (LOC389526), mRNA
 XM_374226 Homo sapiens LOC389533 (LOC389533), mRNA
 XM_374227 Homo sapiens LOC389534 (LOC389534), mRNA
 XM_374232 Homo sapiens LOC389556 (LOC389556), mRNA
 XM_374236 Homo sapiens LOC389562 (LOC389562), mRNA
 XM_374242 Homo sapiens LOC389605 (LOC389605), mRNA
 XM_374243 Homo sapiens LOC389608 (LOC389608), mRNA
 XM_374244 Homo sapiens LOC389609 (LOC389609), mRNA
 XM_374245 Homo sapiens LOC389612 (LOC389612), mRNA
 XM_374247 Homo sapiens LOC389623 (LOC389623), mRNA
 XM_374248 Homo sapiens LOC389624 (LOC389624), mRNA
 XM_374249 Homo sapiens similar to ZNF169 protein (LOC389628), mRNA
 XM_374250 Homo sapiens LOC389629 (LOC389629), mRNA
 XM_374251 Homo sapiens LOC389632 (LOC389632), mRNA
 XM_374252 Homo sapiens LOC389634 (LOC389634), mRNA
 XM_374253 Homo sapiens LOC389635 (LOC389635), mRNA
 XM_374254 Homo sapiens LOC389636 (LOC389636), mRNA
 XM_374255 Homo sapiens LOC389637 (LOC389637), mRNA
 XM_374256 Homo sapiens LOC389638 (LOC389638), mRNA
 XM_374257 Homo sapiens hypothetical protein LOC286114 (LOC286114), mRNA
 XM_374258 Homo sapiens LOC389639 (LOC389639), mRNA
 XM_374259 Homo sapiens LOC389640 (LOC389640), mRNA
 XM_374260 Homo sapiens LOC389641 (LOC389641), mRNA
 XM_374261 Homo sapiens LOC389642 (LOC389642), mRNA
 XM_374262 Homo sapiens LOC389645 (LOC389645), mRNA
 XM_374263 Homo sapiens LOC389647 (LOC389647), mRNA
 XM_374264 Homo sapiens LOC389648 (LOC389648), mRNA
 XM_374266 Homo sapiens LOC389650 (LOC389650), mRNA
 XM_374267 Homo sapiens LOC389654 (LOC389654), mRNA
 XM_374268 Homo sapiens LOC389656 (LOC389656), mRNA
 XM_374269 Homo sapiens LOC389657 (LOC389657), mRNA
 XM_374270 Homo sapiens LOC389659 (LOC389659), mRNA
 XM_374271 Homo sapiens LOC389665 (LOC389665), mRNA
 XM_374272 Homo sapiens LOC389666 (LOC389666), mRNA
 XM_374273 Homo sapiens LOC389669 (LOC389669), mRNA
 XM_374274 Homo sapiens LOC389670 (LOC389670), mRNA
 XM_374275 Homo sapiens LOC389671 (LOC389671), mRNA
 XM_374276 Homo sapiens LOC389676 (LOC389676), mRNA
 XM_374277 Homo sapiens LOC389683 (LOC389683), mRNA
 XM_374278 Homo sapiens LOC389684 (LOC389684), mRNA
 XM_374279 Homo sapiens LOC389685 (LOC389685), mRNA
 XM_374280 Homo sapiens LOC389686 (LOC389686), mRNA
 XM_374281 Homo sapiens LOC389687 (LOC389687), mRNA
 XM_374282 Homo sapiens LOC389688 (LOC389688), mRNA
 XM_374283 Homo sapiens LOC389689 (LOC389689), mRNA

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XM_374284 Homo sapiens LOC389691 (LOC389691), mRNA
XM_374285 Homo sapiens LOC389693 (LOC389693), mRNA
XM_374286 Homo sapiens LOC389695 (LOC389695), mRNA
XM_374287 Homo sapiens LOC389696 (LOC389696), mRNA
XM_374289 Homo sapiens LOC389702 (LOC389702), mRNA
XM_374290 Homo sapiens LOC389703 (LOC389703), mRNA
XM_374291 Homo sapiens LOC389706 (LOC389706), mRNA
XM_374292 Homo sapiens LOC389707 (LOC389707), mRNA
XM_374293 Homo sapiens LOC389718 (LOC389718), mRNA
XM_374294 Homo sapiens LOC389720 (LOC389720), mRNA
XM_374295 Homo sapiens LOC389767 (LOC389767), mRNA
XM_374296 Homo sapiens LOC389771 (LOC389771), mRNA
XM_374297 Homo sapiens LOC389773 (LOC389773), mRNA
XM_374298 Homo sapiens LOC389774 (LOC389774), mRNA
XM_374299 Homo sapiens LOC389775 (LOC389775), mRNA
XM_374300 Homo sapiens LOC389777 (LOC389777), mRNA
XM_374301 Homo sapiens LOC389778 (LOC389778), mRNA
XM_374302 Homo sapiens LOC389782 (LOC389782), mRNA
XM_374303 Homo sapiens LOC389783 (LOC389783), mRNA
XM_374304 Homo sapiens LOC389784 (LOC389784), mRNA
XM_374305 Homo sapiens LOC389793 (LOC389793), mRNA
XM_374306 Homo sapiens LOC389794 (LOC389794), mRNA
XM_374307 Homo sapiens LOC389801 (LOC389801), mRNA
XM_374308 Homo sapiens LOC389804 (LOC389804), mRNA
XM_374309 Homo sapiens LOC389806 (LOC389806), mRNA
XM_374313 Homo sapiens LOC389810 (LOC389810), mRNA
XM_374315 Homo sapiens LOC389815 (LOC389815), mRNA
XM_374317 Homo sapiens LOC389831 (LOC389831), mRNA
XM_374318 Homo sapiens LOC389834 (LOC389834), mRNA
XM_374319 Homo sapiens hypothetical protein LOC339457 (LOC339457), mRNA
XM_374323 Homo sapiens LOC389841 (LOC389841), mRNA
XM_374325 Homo sapiens LOC389846 (LOC389846), mRNA
XM_374326 Homo sapiens LOC389858 (LOC389858), mRNA
XM_374327 Homo sapiens LOC389861 (LOC389861), mRNA
XM_374328 Homo sapiens LOC389863 (LOC389863), mRNA
XM_374329 Homo sapiens LOC389865 (LOC389865), mRNA
XM_374330 Homo sapiens LOC389867 (LOC389867), mRNA
XM_374331 Homo sapiens LOC389868 (LOC389868), mRNA
XM_374333 Homo sapiens LOC389878 (LOC389878), mRNA
XM_374336 Homo sapiens LOC389889 (LOC389889), mRNA
XM_374337 Homo sapiens LOC389890 (LOC389890), mRNA
XM_374338 Homo sapiens LOC389893 (LOC389893), mRNA
XM_374339 Homo sapiens LOC389896 (LOC389896), mRNA
XM_374340 Homo sapiens LOC389897 (LOC389897), mRNA
XM_374341 Homo sapiens LOC389908 (LOC389908), mRNA
XM_374342 Homo sapiens LOC389909 (LOC389909), mRNA
XM_374343 Homo sapiens LOC389910 (LOC389910), mRNA
XM_374349 Homo sapiens LOC389931 (LOC389931), mRNA
XM_374352 Homo sapiens LOC390367 (LOC390367), mRNA
XM_374354 Homo sapiens LOC390550 (LOC390550), mRNA
XM_374355 Homo sapiens LOC390565 (LOC390565), mRNA
XM_374356 Homo sapiens LOC390568 (LOC390568), mRNA
XM_374357 Homo sapiens LOC390749 (LOC390749), mRNA
XM_374358 Homo sapiens LOC390751 (LOC390751), mRNA
XM_374359 Homo sapiens LOC390821 (LOC390821), mRNA
XM_374361 Homo sapiens LOC390870 (LOC390870), mRNA
XM_374362 Homo sapiens LOC390899 (LOC390899), mRNA

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XM_374364 Homo sapiens LOC391067 (LOC391067), mRNA
 XM_374365 Homo sapiens LOC391238 (LOC391238), mRNA
 XM_374367 Homo sapiens similar to hypothetical protein AN2833.2 (LOC391727), mRNA
 XM_374368 Homo sapiens LOC391728 (LOC391728), mRNA
 XM_374369 Homo sapiens LOC391729 (LOC391729), mRNA
 XM_374370 Homo sapiens LOC392160 (LOC392160), mRNA
 XM_374371 Homo sapiens LOC392177 (LOC392177), mRNA
 XM_374372 Homo sapiens LOC392238 (LOC392238), mRNA
 XM_374377 Homo sapiens LOC392580 (LOC392580), mRNA
 XM_374378 Homo sapiens LOC392581 (LOC392581), mRNA
 XM_374379 Homo sapiens LOC392583 (LOC392583), mRNA
 XM_374382 Homo sapiens similar to cAMP-dependent protein kinase type I-beta regulat
 XM_374386 Homo sapiens similar to RIKEN cDNA A930017N06 gene (LOC392617), mR
 XM_374388 Homo sapiens similar to KIAA1691 protein (LOC392619), mRNA
 XM_374389 Homo sapiens KIAA1950 protein (KIAA1950), mRNA
 XM_374396 Homo sapiens ubiquitin specific protease 42 (USP42), mRNA
 XM_374399 Homo sapiens similar to matrilin 2 precursor (LOC392630), mRNA
 XM_374401 Homo sapiens similar to alpha-2 macroglobulin family protein VIP (LOC392
 XM_374404 Homo sapiens KIAA0960 protein (KIAA0960), mRNA
 XM_374405 Homo sapiens similar to Neurogenic locus notch homolog protein 1 precurs
 XM_374406 Homo sapiens similar to RIKEN cDNA A530016O06 gene (LOC392636), mR
 XM_374413 Homo sapiens similar to Neuronal protein 3.1 (p311 protein) (LOC392642), n
 XM_374414 Homo sapiens similar to mKIAA0038 protein (LOC392647), mRNA
 XM_374419 Homo sapiens similar to RP9 protein (LOC392652), mRNA
 XM_374422 Homo sapiens KIAA0877 protein (KIAA0877), mRNA
 XM_374425 Homo sapiens KIAA0895 protein (KIAA0895), mRNA
 XM_374427 Homo sapiens similar to T-cell receptor gamma chain V region PT-gamma-1
 XM_374428 Homo sapiens LOC392661 (LOC392661), mRNA
 XM_374430 Homo sapiens hypothetical protein DKFZp761I2123 (DKFZp761I2123), mR
 XM_374431 Homo sapiens similar to myosin IG (LOC392665), mRNA
 XM_374432 Homo sapiens similar to KIAA0363 (LOC392666), mRNA
 XM_374434 Homo sapiens LOC392668 (LOC392668), mRNA
 XM_374435 Homo sapiens hypothetical protein LOC136288 (LOC136288), mRNA
 XM_374437 Homo sapiens LOC392670 (LOC392670), mRNA
 XM_374439 Homo sapiens similar to LanC lantibiotic synthetase component C-like 2; tesl
 XM_374460 Homo sapiens similar to Williams-Beuren syndrome critical region protein 19
 XM_374461 Homo sapiens similar to PMS5 homolog mismatch repair protein - human (L
 XM_374473 Homo sapiens similar to general transcription factor II, I isoform 1; BTK-asso
 XM_374475 Homo sapiens similar to PMS2L13 (LOC392729), mRNA
 XM_374479 Homo sapiens similar to Williams-Beuren syndrome critical region protein 19
 XM_374481 Homo sapiens similar to dextx 2 (LOC392736), mRNA
 XM_374483 Homo sapiens similar to 60S ribosomal protein L7a (Surfeit locus protein 3) (L
 XM_374484 Homo sapiens similar to Piccolo protein (Aczonin) (LOC392742), mRNA
 XM_374487 Homo sapiens similar to Six transmembrane epithelial antigen of prostate (L
 XM_374489 Homo sapiens hypothetical protein FLJ39885 (FLJ39885), mRNA
 XM_374490 Homo sapiens similar to ribosomal protein S27 (LOC392748), mRNA
 XM_374491 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 9A (PPP
 XM_374496 Homo sapiens similar to ZNF36 protein (LOC392756), mRNA
 XM_374498 Homo sapiens similar to CG14977-PA (LOC392758), mRNA
 XM_374501 Homo sapiens similar to intestinal mucin 3 (LOC392762), mRNA
 XM_374502 Homo sapiens similar to intestinal mucin (LOC392763), mRNA
 XM_374506 Homo sapiens similar to Williams-Beuren syndrome critical region protein 19
 XM_374514 Homo sapiens similar to laminin beta-4 chain precursor (LOC392775), mRN
 XM_374517 Homo sapiens similar to zinc finger protein 312; forebrain embryonic zinc fin
 XM_374518 Homo sapiens similar to cardiac leiomodlin (LOC392780), mRNA
 XM_374519 Homo sapiens similar to 40S ribosomal protein S2 (LOC392781), mRNA
 XM_374522 Homo sapiens similar to Filamin C (Gamma-filamin) (Filamin 2) (Protein FLN

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XM_374526 Homo sapiens similar to eukaryotic translation elongation factor 1 beta 2; eul
 XM_374529 Homo sapiens KIAA1466 protein (KIAA1466), mRNA
 XM_374530 Homo sapiens nucleoporin 205kDa (NUP205), mRNA
 XM_374538 Homo sapiens KIAA1277 (KIAA1277), mRNA
 XM_374579 Homo sapiens hypothetical protein LOC155054 (LOC155054), mRNA
 XM_374581 Homo sapiens similar to KIAA1862 protein (LOC392836), mRNA
 XM_374583 Homo sapiens similar to actin-related protein Arp11 (LOC392838), mRNA
 XM_374584 Homo sapiens similar to amyotrophic lateral sclerosis 2 (juvenile) chromosome
 XM_374585 Homo sapiens similar to hypothetical protein 4931409K22 (LOC392842), mF
 XM_374586 Homo sapiens similar to hypothetical protein FLJ22527 (LOC392843), mRN/
 XM_374589 Homo sapiens LOC392848 (LOC392848), mRNA
 XM_374590 Homo sapiens similar to C7orf3 protein (LOC392850), mRNA
 XM_374591 Homo sapiens LOC392851 (LOC392851), mRNA
 XM_374592 Homo sapiens similar to THAP domain containing 5 (LOC392852), mRNA
 XM_374600 Homo sapiens similar to GluR-delta2 philic-protein (LOC392862), mRNA
 XM_374603 Homo sapiens similar to cell division cycle 10 homolog (LOC392884), mRNA
 XM_374604 Homo sapiens similar to T-cell receptor (V-J-C) precursor (LOC392887), mR
 XM_374605 Homo sapiens similar to T-cell receptor gamma chain V region PT-gamma-1,
 XM_374607 Homo sapiens similar to Splicing factor, arginine/serine-rich, 46kD (LOC3928
 XM_374608 Homo sapiens hypothetical protein MGC26484 (MGC26484), mRNA
 XM_374625 Homo sapiens similar to hypothetical protein (LOC392943), mRNA
 XM_374627 Homo sapiens guanine nucleotide binding protein, alpha transducing 3 (GNA
 XM_374628 Homo sapiens similar to dynein, cytoplasmic, light peptide; 8kD LC; dynein L
 XM_374636 Homo sapiens similar to acyl CoA:monacylglycerol acyltransferase 2 (LOC3
 XM_374637 Homo sapiens similar to alpha-2-glycoprotein 1, zinc; Alpha-2-glycoprotein, z
 XM_374638 Homo sapiens similar to Cohesin subunit SA-3 (Stromal antigen 3) (SCC3 hc
 XM_374646 Homo sapiens similar to ribosomal protein L18; 60S ribosomal protein L18 (L
 XM_374648 Homo sapiens similar to RIKEN cDNA 1500011L16 (LOC392982), mRNA
 XM_374650 Homo sapiens similar to ribosomal protein S14 (LOC392988), mRNA
 XM_374653 Homo sapiens similar to CG6293-PA (LOC392993), mRNA
 XM_374655 Homo sapiens similar to dJ753D5.2 (novel protein similar to RPS17 (40S rib
 XM_374657 Homo sapiens similar to dGTPase (EC 3.1.5.1) - mouse (fragment) (LOC393
 XM_374680 Homo sapiens similar to seven transmembrane helix receptor (LOC393037),
 XM_374682 Homo sapiens similar to Olfactory receptor 6B1 (Olfactory receptor 7-3) (OR;
 XM_374683 Homo sapiens similar to olfactory receptor MOR261-13 (LOC393046), mRNA/
 XM_374684 Homo sapiens similar to olfactory receptor MOR261-1 (LOC393047), mRNA
 XM_374687 Homo sapiens similar to seven transmembrane helix receptor (LOC393051),
 XM_374689 Homo sapiens similar to seven transmembrane helix receptor (LOC393053),
 XM_374694 Homo sapiens similar to Mtr3 (mRNA transport regulator 3)-homolog; Mtr3 (n
 XM_374699 Homo sapiens similar to gastrulation brain homeobox 1 (LOC393070), mRN/
 XM_374702 Homo sapiens similar to envelope protein (LOC393073), mRNA
 XM_374705 Homo sapiens similar to polycystic kidney disease 1-like 3 (LOC393078), mF
 XM_374706 Homo sapiens similar to 60S RIBOSOMAL PROTEIN L21 (LOC393079), mR
 XM_374710 Homo sapiens LOC392620 (LOC392620), mRNA
 XM_374711 Homo sapiens LOC392621 (LOC392621), mRNA
 XM_374713 Homo sapiens similar to Zinc finger protein 268 (Zinc finger protein HZF3) (L
 XM_374715 Homo sapiens LOC392641 (LOC392641), mRNA
 XM_374721 Homo sapiens LOC392657 (LOC392657), mRNA
 XM_374722 Homo sapiens LOC392659 (LOC392659), mRNA
 XM_374730 Homo sapiens LOC392702 (LOC392702), mRNA
 XM_374734 Homo sapiens LOC392726 (LOC392726), mRNA
 XM_374735 Homo sapiens LOC392730 (LOC392730), mRNA
 XM_374741 Homo sapiens LOC392749 (LOC392749), mRNA
 XM_374751 Homo sapiens LOC392790 (LOC392790), mRNA
 XM_374752 Homo sapiens LOC392791 (LOC392791), mRNA
 XM_374761 Homo sapiens LOC392849 (LOC392849), mRNA
 XM_374763 Homo sapiens LOC393076 (LOC393076), mRNA

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XM_374764 Homo sapiens LOC393080 (LOC393080), mRNA
 XM_374765 Homo sapiens similar to cDNA sequence BC016423 (LOC399712), mRNA
 XM_374766 Homo sapiens hypothetical gene supported by AK128185 (LOC399715), mR
 XM_374767 Homo sapiens similar to hypothetical protein (LOC399716), mRNA
 XM_374768 Homo sapiens USP6 N-terminal like (USP6NL), mRNA
 XM_374769 Homo sapiens similar to USP6NL protein (LOC399718), mRNA
 XM_374770 Homo sapiens similar to seven transmembrane helix receptor (LOC399719),
 XM_374779 Homo sapiens ankyrin repeat domain 26 (ANKRD26), mRNA
 XM_374781 Homo sapiens hypothetical protein LOC220906 (LOC220906), mRNA
 XM_374782 Homo sapiens similar to supervillin isoform 2; membrane-associated F-actin
 XM_374786 Homo sapiens similar to hypothetical protein LOC349114 (LOC399744), mR
 XM_374787 Homo sapiens similar to hypothetical protein FLJ40432 (LOC399747), mRN/
 XM_374792 Homo sapiens similar to Hypothetical protein KIAA0514 (LOC399755), mRN/
 XM_374799 Homo sapiens similar to ARF GTPase-activating protein (LOC399758), mRN
 XM_374801 Homo sapiens similar to ARF GTPase-activating protein (LOC399761), mRN
 XM_374802 Homo sapiens similar to DKFZP566K0524 protein (LOC399762), mRNA
 XM_374803 Homo sapiens similar to LINE-1 REVERSE TRANSCRIPTASE HOMOLOG (/
 XM_374807 Homo sapiens similar to ARF GTPase-activating protein (LOC399769), mRN
 XM_374809 Homo sapiens similar to activator of S phase kinase (LOC399777), mRNA
 XM_374810 Homo sapiens similar to activator of S phase kinase (LOC399778), mRNA
 XM_374813 Homo sapiens similar to peptidylprolyl isomerase A (cyclophilin A) (LOC3997
 XM_374817 Homo sapiens similar to bA182L21.1 (novel protein similar to hypothetical pr
 XM_374829 Homo sapiens programmed cell death 11 (PCDD11), mRNA
 XM_374830 Homo sapiens similar to hypothetical protein LOC119395 (LOC399807), mR
 XM_374831 Homo sapiens SH3 multiple domains 1 (SH3MD1), mRNA
 XM_374832 Homo sapiens similar to ribosomal protein L13a; 60S ribosomal protein L13a
 XM_374835 Homo sapiens similar to RIKEN cDNA 1700022C21 (LOC399814), mRNA
 XM_374836 Homo sapiens similar to hypothetical protein FLJ13490 (LOC399815), mRN/
 XM_374839 Homo sapiens similar to CG9643-PA (LOC399818), mRNA
 XM_374840 Homo sapiens similar to CG15021-PA (LOC399819), mRNA
 XM_374842 Homo sapiens hypothetical gene supported by AK094354 (LOC399821), mR
 XM_374844 Homo sapiens similar to hypothetical protein B130055A05 (LOC399823), mF
 XM_374847 Homo sapiens similar to shadow of prion protein; Shadoo (LOC399831), mR
 XM_374851 Homo sapiens similar to Synaptotagmin XV (SytXV) (Chr10Syt) (LOC399837
 XM_374852 Homo sapiens similar to double homeobox protein (LOC399839), mRNA
 XM_374854 Homo sapiens similar to nuclear receptor coactivator 4; RET-activating gene
 XM_374855 Homo sapiens similar to Glutamate dehydrogenase 1, mitochondrial precursor
 XM_374857 Homo sapiens similar to bA476I15.3 (novel protein similar to septin) (LOC39/
 XM_374858 Homo sapiens hypothetical gene supported by AK093729; AK128780; AL11;
 XM_374860 Homo sapiens tumor protein p53 Inducible protein 5 (TP53I5), mRNA
 XM_374864 Homo sapiens hypothetical gene supported by AK091259 (LOC399857), mR
 XM_374873 Homo sapiens similar to RIKEN cDNA 3830422K02 (LOC399870), mRNA
 XM_374877 Homo sapiens hypothetical protein DKFZp779M0652 (DKFZp779M0652), ml
 XM_374879 Homo sapiens hypothetical protein LOC114971 (LOC114971), mRNA
 XM_374880 Homo sapiens hypothetical gene supported by BC065704 (LOC399888), mR
 XM_374882 Homo sapiens similar to Metabotropic glutamate receptor 5 precursor (mGlu
 XM_374885 Homo sapiens hypothetical gene supported by AK128188 (LOC399898), mR
 XM_374890 Homo sapiens hypothetical gene supported by AK093779 (LOC399900), mR
 XM_374893 Homo sapiens LOC399904 (LOC399904), mRNA
 XM_374896 Homo sapiens similar to pecanex-like 3 (LOC399909), mRNA
 XM_374899 Homo sapiens similar to Gag-Pro-Pol protein (LOC399913), mRNA
 XM_374900 Homo sapiens similar to beta-1,4-mannosyltransferase; beta-1,4 mannosyltr
 XM_374902 Homo sapiens similar to polymerase (LOC399917), mRNA
 XM_374904 Homo sapiens similar to SH3 and multiple ankyrin repeat domains protein 2 i
 XM_374905 Homo sapiens similar to proline rich synapse associated protein 1 isoform E;
 XM_374906 Homo sapiens hypothetical gene supported by AK124096 (LOC399923), mR
 XM_374907 Homo sapiens SH3 multiple domains 3 (SH3MD3), mRNA

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XM_374909 Homo sapiens similar to hypothetical protein (LOC399925), mRNA
XM_374911 Homo sapiens similar to ribosomal protein S12 (LOC399927), mRNA
XM_374912 Homo sapiens X-ray radiation resistance associated 1 (XRR1), mRNA
XM_374915 Homo sapiens hypothetical protein LOC283219 (LOC283219), mRNA
XM_374917 Homo sapiens similar to tripartite motif-containing 51 (LOC399937), mRNA
XM_374919 Homo sapiens similar to tripartite motif protein 48; TRIM48 (LOC399939), mRNA
XM_374920 Homo sapiens similar to tripartite motif-containing 51 (LOC399940), mRNA
XM_374922 Homo sapiens KIAA1731 protein (KIAA1731), mRNA
XM_374927 Homo sapiens KIAA1377 protein (KIAA1377), mRNA
XM_374930 Homo sapiens similar to expressed sequence A1593442 (LOC399947), mRNA
XM_374932 Homo sapiens hypothetical gene supported by AB096245 (LOC399948), mRNA
XM_374933 Homo sapiens similar to RIKEN cDNA 4833427G06 (LOC399949), mRNA
XM_374936 Homo sapiens KIAA1052 protein (KIAA1052), mRNA
XM_374937 Homo sapiens hypothetical gene supported by AK127233 (LOC399957), mRNA
XM_374944 Homo sapiens similar to LVLF3112 (LOC399967), mRNA
XM_374945 Homo sapiens similar to Seminal vesicle protein 7 precursor (SVS VII) (Caltri
XM_374948 Homo sapiens similar to LINE-1 REVERSE TRANSCRIPTASE HOMOLOG (L
XM_374949 Homo sapiens similar to Sorting nexin 19 (LOC399979), mRNA
XM_374952 Homo sapiens hypothetical gene supported by AK126822 (LOC399990), mRNA
XM_374959 Homo sapiens similar to DDX11 protein (LOC399999), mRNA
XM_374965 Homo sapiens similar to BC004636 protein (LOC400010), mRNA
XM_374967 Homo sapiens similar to carboxy-terminal modulator protein isoform a (LOC
XM_374972 Homo sapiens similar to MUC19 (LOC400023), mRNA
XM_374973 Homo sapiens similar to hypothetical protein (L1H 3 region) - human (LOC4C
XM_374976 Homo sapiens hypothetical protein LOC283331 (LOC283331), mRNA
XM_374981 Homo sapiens similar to DKFZP564K247 protein (LOC400038), mRNA
XM_374982 Homo sapiens similar to oriLyt TD-element binding protein 7 (LOC400040), mRNA
XM_374983 Homo sapiens KIAA0748 gene product (KIAA0748), mRNA
XM_374985 Homo sapiens KIAA0352 gene product (KIAA0352), mRNA
XM_374987 Homo sapiens similar to 60S ribosomal protein L26 (LOC400055), mRNA
XM_374989 Homo sapiens KIAA0373 gene product (KIAA0373), mRNA
XM_374995 Homo sapiens similar to ribosomal protein L18a; 60S ribosomal protein L18a
XM_374996 Homo sapiens AMP-activated protein kinase family member 5 (ARK5), mRNA
XM_374997 Homo sapiens hypothetical gene supported by AK097461; BC046185 (LOC4
XM_374999 Homo sapiens hypothetical gene supported by AK124947 (LOC400077), mRNA
XM_375000 Homo sapiens KIAA1853 protein (KIAA1853), mRNA
XM_375004 Homo sapiens similar to Dynein heavy chain at 89D CG1842-PA (LOC40008
XM_375005 Homo sapiens similar to Dynein heavy chain at 89D CG1842-PA (LOC40008
XM_375007 Homo sapiens hypothetical gene supported by AK123815 (LOC400093), mRNA
XM_375013 Homo sapiens hypothetical gene supported by AK127292; AK128225 (LOC4
XM_375018 Homo sapiens hypothetical gene supported by AK092066 (LOC400121), mRNA
XM_375023 Homo sapiens hypothetical gene supported by AJ412041 (LOC400132), mRNA
XM_375027 Homo sapiens hypothetical protein LOC283491 (LOC283491), mRNA
XM_375029 Homo sapiens hypothetical protein LOC338862 (LOC338862), mRNA
XM_375031 Homo sapiens hypothetical gene supported by AK093158 (LOC400145), mRNA
XM_375032 Homo sapiens TBC1 domain family, member 4 (TBC1D4), mRNA
XM_375033 Homo sapiens hypothetical protein LOC144776 (LOC144776), mRNA
XM_375035 Homo sapiens similar to 40S ribosomal protein S26 (LOC400156), mRNA
XM_375038 Homo sapiens hypothetical gene supported by AK129953 (LOC400165), mRNA
XM_375039 Homo sapiens similar to LRRGT00052 (LOC400169), mRNA
XM_375041 Homo sapiens similar to CLL-associated antigen KW-1 splice variant 1 (LOC
XM_375042 Homo sapiens DKFZP434B061 protein (DKFZP434B061), mRNA
XM_375045 Homo sapiens chromosome 14 open reading frame 92 (C14orf92), mRNA
XM_375065 Homo sapiens zinc finger protein 409 (ZNF409), mRNA
XM_375067 Homo sapiens similar to peroxisomal short-chain alcohol dehydrogenase; N/A
XM_375074 Homo sapiens KIAA0391 (KIAA0391), mRNA
XM_375075 Homo sapiens similar to STELLA (LOC400206), mRNA

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XM_375076 Homo sapiens hypothetical gene supported by AK124214 (LOC400207), mR
 XM_375077 Homo sapiens chromosome 14 open reading frame 25 (C14orf25), mRNA
 XM_375078 Homo sapiens similar to small acidic protein; small acidic protein sid2057p (L
 XM_375080 Homo sapiens KIAA0831 (KIAA0831), mRNA
 XM_375081 Homo sapiens hypothetical gene supported by BX248296 (LOC400214), mR
 XM_375084 Homo sapiens similar to hypothetical protein (LOC400219), mRNA
 XM_375085 Homo sapiens KIAA1393 (KIAA1393), mRNA
 XM_375086 Homo sapiens zinc finger and BTB domain containing 1 (ZBTB1), mRNA
 XM_375087 Homo sapiens pleckstrin homology domain containing, family H (with MyTH4
 XM_375088 Homo sapiens hypothetical gene supported by AK097098 (LOC400223), mR
 XM_375090 Homo sapiens similar to pleckstrin homology domain protein (5V327) (LOC4
 XM_375099 Homo sapiens hypothetical protein LOC283585 (LOC283585), mRNA
 XM_375101 Homo sapiens similar to RIKEN cDNA 061001D24 (LOC400239), mRNA
 XM_375105 Homo sapiens KIAA0329 (KIAA0329), mRNA
 XM_375108 Homo sapiens similar to RIKEN cDNA A530016L24 gene (LOC400258), mR
 XM_375145 Homo sapiens similar to breast cancer antigen NY-BR-1 (LOC400296), mRN
 XM_375147 Homo sapiens similar to breast cancer anti-estrogen resistance 1; Crk-assoc
 XM_375148 Homo sapiens similar to Ribosome biogenesis protein BMS1 homolog (LOC
 XM_375150 Homo sapiens similar to protein kinase CHK2 isoform b; checkpoint-like prot
 XM_375152 Homo sapiens similar to hypothetical protein (LOC400304), mRNA
 XM_375153 Homo sapiens hypothetical protein DKFZp547L112 (DKFZP547L112), mRN/
 XM_375154 Homo sapiens similar to KIAA0125 (LOC400309), mRNA
 XM_375157 Homo sapiens similar to Ribosome biogenesis protein BMS1 homolog (LOC
 XM_375163 Homo sapiens similar to hypothetical protein FLJ38144 (LOC400320), mRN/
 XM_375165 Homo sapiens hypothetical protein DKFZp434P162 (LOC390535), mRNA
 XM_375170 Homo sapiens similar to KIAA1971 protein (LOC400336), mRNA
 XM_375171 Homo sapiens DKFZP434L187 protein (DKFZP434L187), mRNA
 XM_375174 Homo sapiens hypothetical gene supported by BC037839 (LOC400340), mR
 XM_375176 Homo sapiens similar to RIKEN cDNA 653040I14 gene (LOC400342), mR
 XM_375178 Homo sapiens similar to hect domain and RLD 2 (LOC400344), mRNA
 XM_375179 Homo sapiens similar to hect domain and RLD 2 (LOC400345), mRNA
 XM_375181 Homo sapiens KIAA1018 protein (KIAA1018), mRNA
 XM_375183 Homo sapiens hypothetical gene supported by BC037839 (LOC400353), mR
 XM_375185 Homo sapiens formin (limb deformity) (FMN), mRNA
 XM_375187 Homo sapiens nuclear protein in testis (NUT), mRNA
 XM_375190 Homo sapiens hypothetical gene supported by AK093014 (LOC400359), mR
 XM_375191 Homo sapiens hypothetical gene supported by BX647708 (LOC400360), mR
 XM_375196 Homo sapiens similar to KIAA0377 gene product (LOC400367), mRNA
 XM_375200 Homo sapiens similar to ubiquitin specific proteinase 50 (LOC400372), mRN
 XM_375203 Homo sapiens hypothetical gene supported by AK126787 (LOC400376), mR
 XM_375207 Homo sapiens similar to RIKEN cDNA 1110004B15 (LOC400380), mRNA
 XM_375209 Homo sapiens likely ortholog of mouse klotho lactase-phlorizin hydrolase reli
 XM_375210 Homo sapiens similar to zinc finger and BTB domain containing 8; BTB/POZ
 XM_375224 Homo sapiens similar to cervical cancer suppressor-1 (LOC400410), mRNA
 XM_375226 Homo sapiens similar to FLJ40113 protein (LOC400414), mRNA
 XM_375228 Homo sapiens similar to RIKEN cDNA 3010021M21 (LOC400416), mRNA
 XM_375230 Homo sapiens similar to KIAA1920 protein (LOC400418), mRNA
 XM_375235 Homo sapiens similar to hypothetical protein FLJ22795 (LOC400423), mRN/
 XM_375239 Homo sapiens similar to KIAA1920 protein (LOC400428), mRNA
 XM_375240 Homo sapiens similar to hypothetical protein FLJ22795 (LOC400429), mRN/
 XM_375242 Homo sapiens similar to golgi autoantigen golgin subfamily a2-like (LOC400
 XM_375243 Homo sapiens KIAA1920 protein (KIAA1920), mRNA
 XM_375246 Homo sapiens similar to hypothetical protein DKFZp434I1020 (LOC400435),
 XM_375247 Homo sapiens KIAA0211 gene product (KIAA0211), mRNA
 XM_375248 Homo sapiens similar to KIAA1920 protein (LOC400436), mRNA
 XM_375249 Homo sapiens similar to hypothetical protein FLJ22795 (LOC400437), mRN/
 XM_375252 Homo sapiens LOC400442 (LOC400442), mRNA

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XM_375260 Homo sapiens hypothetical gene supported by AK075564; BC060873 (LOC4
 XM_375261 Homo sapiens similar to Nonhistone chromosomal protein HMG-14 (High-mc
 XM_375263 Homo sapiens similar to IFM9370 (LOC400454), mRNA
 XM_375266 Homo sapiens LOC400461 (LOC400461), mRNA
 XM_375268 Homo sapiens similar to RIKEN cDNA 3010021M21 (LOC400464), mRNA
 XM_375269 Homo sapiens similar to Dynamin-1 (D100) (Dynamin, brain) (B-dynamin) (L
 XM_375272 Homo sapiens similar to hypothetical protein FLJ36144 (LOC400468), mRNA
 XM_375274 Homo sapiens similar to solute carrier family 22 member 4; organic cation tr
 XM_375275 Homo sapiens similar to hypothetical protein FLJ22795 (LOC400472), mRNA
 XM_375276 Homo sapiens similar to KIAA1920 protein (LOC400473), mRNA
 XM_375277 Homo sapiens similar to FLJ40113 protein (LOC400477), mRNA
 XM_375280 Homo sapiens similar to KIAA1920 protein (LOC400478), mRNA
 XM_375282 Homo sapiens similar to DDX11 protein (LOC400479), mRNA
 XM_375284 Homo sapiens similar to interleukin 9 receptor (LOC400481), mRNA
 XM_375288 Homo sapiens hypothetical protein MGC24381 (MGC24381), mRNA
 XM_375292 Homo sapiens similar to hypothetical protein (LOC400492), mRNA
 XM_375298 Homo sapiens KIAA1987 protein (KIAA1987), mRNA
 XM_375302 Homo sapiens hypothetical gene supported by AK126539 (LOC400499), mR
 XM_375305 Homo sapiens similar to TSG118.1 protein (LOC400506), mRNA
 XM_375306 Homo sapiens similar to hypothetical protein FLJ20581 (LOC400507), mRN
 XM_375307 Homo sapiens similar to hypothetical protein (LOC400508), mRNA
 XM_375308 Homo sapiens similar to FLJ12363 protein (LOC400509), mRNA
 XM_375313 Homo sapiens hypothetical protein MGC9515 (MGC9515), mRNA
 XM_375316 Homo sapiens hypothetical protein LOC283887 (LOC283887), mRNA
 XM_375319 Homo sapiens similar to nuclear pore complex interacting protein (LOC4005
 XM_375320 Homo sapiens similar to apolipoprotein B48 receptor (LOC400514), mRNA
 XM_375325 Homo sapiens similar to nuclear pore complex interacting protein (LOC4005
 XM_375330 Homo sapiens similar to MGC9515 protein (LOC400520), mRNA
 XM_375331 Homo sapiens similar to PI-3-kinase-related kinase SMG-1 isoform 1; lambd
 XM_375333 Homo sapiens hypothetical protein FLJ25404 (FLJ25404), mRNA
 XM_375334 Homo sapiens hypothetical protein LOC283901 (LOC283901), mRNA
 XM_375341 Homo sapiens similar to Ig heavy chain - human (fragment) (LOC400524), m
 XM_375344 Homo sapiens similar to hypothetical protein (LOC400526), mRNA
 XM_375349 Homo sapiens similar to protein phosphatase 2A 48 kDa regulatory subunit i
 XM_375351 Homo sapiens similar to KIAA1501 protein (LOC400529), mRNA
 XM_375352 Homo sapiens similar to KIAA1501 protein (LOC400530), mRNA
 XM_375353 Homo sapiens hypothetical protein LOC146481 (LOC146481), mRNA
 XM_375355 Homo sapiens Nedd4 binding protein 1 (NBP1), mRNA
 XM_375357 Homo sapiens similar to hypothetical protein (LOC400537), mRNA
 XM_375358 Homo sapiens hypothetical protein FLJ25339 (FLJ25339), mRNA
 XM_375359 Homo sapiens brain expressed, associated with Nedd4 (BEAN), mRNA
 XM_375360 Homo sapiens hypothetical gene supported by AK130753 (LOC400539), mR
 XM_375362 Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+ sys
 XM_375363 Homo sapiens similar to PI-3-kinase-related kinase SMG-1 isoform 2; lambd
 XM_375364 Homo sapiens similar to PI-3-kinase-related kinase SMG-1 isoform 2; lambd
 XM_375369 Homo sapiens similar to LHPE306 (LOC400546), mRNA
 XM_375373 Homo sapiens similar to LOC93426 protein (LOC400547), mRNA
 XM_375375 Homo sapiens KIAA0431 protein (KIAA0431), mRNA
 XM_375376 Homo sapiens KIAA0703 gene product (KIAA0703), mRNA
 XM_375377 Homo sapiens KIAA0513 gene product (KIAA0513), mRNA
 XM_375378 Homo sapiens similar to hypothetical protein (LOC400549), mRNA
 XM_375379 Homo sapiens hypothetical gene supported by AK127438 (LOC400555), mR
 XM_375383 Homo sapiens hypothetical gene supported by AK126695 (LOC400559), mR
 XM_375384 Homo sapiens similar to AFG3(ATPase family gene 3)-like 1 (LOC400563), r
 XM_375386 Homo sapiens similar to bA47615.3 (novel protein similar to septin) (LOC40
 XM_375387 Homo sapiens hypothetical gene supported by AK128660 (LOC400566), mR
 XM_375392 Homo sapiens similar to HSPC296 (LOC400569), mRNA

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XM_375397 Homo sapiens KIAA0753 gene product (KIAA0753), mRNA
 XM_375399 Homo sapiens similar to DNA segment, Chr 11, Brigham & Womens Genetic
 XM_375404 Homo sapiens hypothetical protein LOC146850 (LOC146850), mRNA
 XM_375408 Homo sapiens KIAA0672 gene product (KIAA0672), mRNA
 XM_375410 Homo sapiens hypothetical gene supported by AK123100 (LOC400574), mR
 XM_375412 Homo sapiens hypothetical gene supported by AK127731 (LOC400576), mR
 XM_375418 Homo sapiens similar to GRB2-related adaptor protein (LOC400581), mRNA
 XM_375423 Homo sapiens similar to RIKEN cDNA 0610013E23 (LOC400585), mRNA
 XM_375424 Homo sapiens similar to stearyl-CoA desaturase; acyl-CoA desaturase, fatt
 XM_375426 Homo sapiens similar to 60S ribosomal protein L21 (LOC400587), mRNA
 XM_375430 Homo sapiens hypothetical protein LOC201229 (LOC201229), mRNA
 XM_375434 Homo sapiens similar to Very hypothetical protein (LOC400590), mRNA
 XM_375436 Homo sapiens hypothetical gene supported by AK126768 (LOC400591), mR
 XM_375438 Homo sapiens similar to Hypothetical protein KIAA0563 (LOC400594), mRN
 XM_375439 Homo sapiens similar to TBC1 domain family member 3 (Rab GTPase-activ
 XM_375443 Homo sapiens hypothetical protein LOC284100 (LOC284100), mRNA
 XM_375446 Homo sapiens hypothetical protein FLJ11822 (FLJ11822), mRNA
 XM_375449 Homo sapiens hypothetical protein LOC284106 (LOC284106), mRNA
 XM_375452 Homo sapiens similar to keratin associated protein 9.2 (LOC400596), mRNA
 XM_375453 Homo sapiens similar to RIKEN cDNA B830010L13 (LOC400597), mR
 XM_375456 Homo sapiens hypothetical protein DKFzp761G2113 (DKFzp761G2113), mf
 XM_375469 Homo sapiens ProSAPIP2 protein (ProSAPIP2), mRNA
 XM_375471 Homo sapiens KIAA0924 protein (KIAA0924), mRNA
 XM_375475 Homo sapiens hypothetical gene supported by AK126318 (LOC400608), mR
 XM_375478 Homo sapiens similar to RIKEN cDNA 1100001G20 (LOC400610), mRNA
 XM_375482 Homo sapiens similar to U5 snRNP-specific protein, 200 kDa; U5 snRNP-spe
 XM_375484 Homo sapiens similar to adapter protein 162 (LOC400615), mRNA
 XM_375485 Homo sapiens helicase with zinc finger domain (HELZ), mRNA
 XM_375491 Homo sapiens similar to FTO protein (LOC400622), mRNA
 XM_375492 Homo sapiens similar to Ammd protein (LOC400625), mRNA
 XM_375494 Homo sapiens hypothetical gene supported by AK127919 (LOC400627), mR
 XM_375495 Homo sapiens apoptosis-associated tyrosine kinase (AATK), mRNA
 XM_375496 Homo sapiens similar to FLJ00403 protein (LOC400628), mRNA
 XM_375500 Homo sapiens similar to RIKEN cDNA 3110023B02 (MGC16597), mRNA
 XM_375502 Homo sapiens similar to RIKEN cDNA 4921530G04 (LOC400629), mRNA
 XM_375511 Homo sapiens similar to TBC1 domain family member 3 (Rab GTPase-activ
 XM_375514 Homo sapiens similar to pyrroline-5-carboxylate reductase 1 isoform 2; P5C
 XM_375516 Homo sapiens hypothetical protein LOC284121 (LOC284121), mRNA
 XM_375517 Homo sapiens similar to prollyl 4-hydroxylase, beta subunit; v-erb-a avian ery
 XM_375520 Homo sapiens similar to HSPC214 (LOC400637), mRNA
 XM_375527 Homo sapiens hypothetical protein LOC339290 (LOC339290), mRNA
 XM_375537 Homo sapiens similar to thopurine methyltransferase (LOC400650), mRNA
 XM_375543 Homo sapiens similar to 40S ribosomal protein S3a (LOC400652), mRNA
 XM_375544 Homo sapiens hypothetical gene supported by AK126293 (LOC400658), mR
 XM_375545 Homo sapiens hypothetical gene supported by AK126829 (LOC400661), mR
 XM_375548 Homo sapiens similar to bA476115.3 (novel protein similar to septin) (LOC40
 XM_375549 Homo sapiens similar to hypothetical protein LOC349114 (LOC400664), mR
 XM_375550 Homo sapiens similar to Protein Sck (LOC400665), mRNA
 XM_375551 Homo sapiens hypothetical gene supported by AK127589 (LOC400666), mR
 XM_375552 Homo sapiens similar to GLGL782 (LOC400668), mRNA
 XM_375553 Homo sapiens KIAA0963 (KIAA0963), mRNA
 XM_375557 Homo sapiens NY-REN-24 antigen (NY-REN-24), mRNA
 XM_375558 Homo sapiens KIAA1881 (KIAA1881), mRNA
 XM_375559 Homo sapiens scaffold attachment factor B2 (SAFB2), mRNA
 XM_375560 Homo sapiens similar to expressed sequence A1662250 (LOC400673), mRN
 XM_375563 Homo sapiens hypothetical protein FLJ38149 (FLJ38149), mRNA
 XM_375568 Homo sapiens ZFP-36 for a zinc finger protein (HSZFP36), mRNA

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XM_375569 Homo sapiens hypothetical protein DKFZp43411610 (DKFZp43411610), mRNA
 XM_375575 Homo sapiens similar to hypothetical protein (LOC400677), mRNA
 XM_375583 Homo sapiens hypothetical gene supported by AK128109 (LOC400679), mRNA
 XM_375589 Homo sapiens hypothetical gene supported by BC041864 (LOC400681), mRNA
 XM_375590 Homo sapiens similar to zinc finger protein 100; zinc finger protein 100 (Y1) (LOC400682), mRNA
 XM_375593 Homo sapiens zinc finger protein 536 (ZNF536), mRNA
 XM_375594 Homo sapiens zinc finger protein 507 (ZNF507), mRNA
 XM_375599 Homo sapiens hypothetical protein FLJ21369 (FLJ21369), mRNA
 XM_375602 Homo sapiens hypothetical gene supported by AK055260 (LOC400687), mRNA
 XM_375603 Homo sapiens similar to comment for location 3447-3655 BLASTX g|110329f|
 XM_375604 Homo sapiens similar to Hypothetical zinc finger protein KIAA1559 (LOC400689), mRNA
 XM_375606 Homo sapiens hypothetical protein DKFZp779O175 (DKFZp779O175), mRNA
 XM_375608 Homo sapiens similar to hypothetical protein (LOC400692), mRNA
 XM_375609 Homo sapiens similar to Hypothetical zinc finger protein KIAA0961 (LOC400693), mRNA
 XM_375614 Homo sapiens similar to Placental protein 13-like (Charcot-Leyden crystal protein) (LOC400694), mRNA
 XM_375618 Homo sapiens similar to pregnancy-specific beta-1 glycoprotein C1 - human (LOC400695), mRNA
 XM_375619 Homo sapiens similar to polycythemia rubra vera 1; cell surface receptor (LO
 XM_375629 Homo sapiens hypothetical gene DKFZp434J0226 (DKFZp434J0226), mRNA
 XM_375631 Homo sapiens hypothetical gene supported by AK124070 (LOC400707), mRNA
 XM_375632 Homo sapiens similar to Serine/threonine protein phosphatase 5 (PP5) (Prot
 XM_375633 Homo sapiens solute carrier family 8 (sodium-calcium exchanger), member 2
 XM_375634 Homo sapiens similar to sialic acid binding Ig-like lectin 11; sialic acid-binding
 XM_375638 Homo sapiens similar to zinc finger protein 534; KRAB domain only 3 (LOC4
 XM_375639 Homo sapiens similar to zinc finger protein KR-ZNF1 (LOC400714), mRNA
 XM_375640 Homo sapiens similar to hypothetical protein MGC48625 (LOC400715), mRNA
 XM_375646 Homo sapiens zinc finger protein 525 (ZNF525), mRNA
 XM_375651 Homo sapiens KIAA1115 (KIAA1115), mRNA
 XM_375654 Homo sapiens hypothetical protein FLJ35258 (FLJ35258), mRNA
 XM_375655 Homo sapiens similar to hypothetical protein A430110N23 (LOC400717), mRNA
 XM_375656 Homo sapiens similar to hypothetical protein A430110N23 (LOC400718), mRNA
 XM_375658 Homo sapiens hypothetical gene supported by AK123294 (LOC400719), mRNA
 XM_375660 Homo sapiens zinc finger protein 264 (ZNF264), mRNA
 XM_375663 Homo sapiens similar to hypothetical protein FLJ23506 (LOC400720), mRNA
 XM_375664 Homo sapiens similar to KIAA2003 protein (LOC400721), mRNA
 XM_375665 Homo sapiens hypothetical protein BC012365 (LOC116412), mRNA
 XM_375667 Homo sapiens similar to bA476115.3 (novel protein similar to septin) (LOC40
 XM_375668 Homo sapiens hypothetical gene supported by AK093729; AK128780; BX64
 XM_375669 Homo sapiens similar to 80S ribosomal protein L23a (LOC400725), mRNA
 XM_375670 Homo sapiens hypothetical gene supported by AK093729; BC062355; BX64
 XM_375671 Homo sapiens similar to bA476115.3 (novel protein similar to septin) (LOC40
 XM_375678 Homo sapiens similar to KIAA1751 protein (LOC400731), mRNA
 XM_375681 Homo sapiens KIAA0495 (KIAA0495), mRNA
 XM_375682 Homo sapiens glycine-, glutamate-, thienylcyclohexylpiperidine-binding prote
 XM_375684 Homo sapiens halcy and enhancer of split (Drosophila) homolog 2 (HES2), m
 XM_375685 Homo sapiens KIAA0469 gene product (KIAA0469), mRNA
 XM_375687 Homo sapiens similar to RIKEN cDNA F730108M23 gene (LOC400734), mRNA
 XM_375688 Homo sapiens similar to hypothetical protein (LOC400736), mRNA
 XM_375690 Homo sapiens similar to RIKEN cDNA 9030409G11 (LOC400737), mRNA
 XM_375695 Homo sapiens hypothetical protein LOC126917 (LOC126917), mRNA
 XM_375696 Homo sapiens similar to ribosomal protein S14 (LOC400744), mRNA
 XM_375697 Homo sapiens KIAA0459 protein (KIAA0459), mRNA
 XM_375698 Homo sapiens hypothetical gene supported by AK124869 (LOC400745), mRNA
 XM_375700 Homo sapiens similar to Hypothetical protein BC005730 (LOC400746), mRNA
 XM_375707 Homo sapiens hypothetical gene supported by AK054768 (LOC400747), mRNA
 XM_375712 Homo sapiens syndecan 3 (N-syndecan) (SDC3), mRNA
 XM_375713 Homo sapiens hypothetical protein LOC284551 (LOC284551), mRNA
 XM_375714 Homo sapiens similar to RIKEN cDNA 1700025K23 (LOC400749), mRNA

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XM_375718 Homo sapiens inositol polyphosphate-5-phosphatase, 75kDa (INPP5B), mRNA
 XM_375720 Homo sapiens regulating synaptic membrane exocytosis 3 (RIMS3), mRNA
 XM_375726 Homo sapiens KIAA0494 gene product (KIAA0494), mRNA
 XM_375729 Homo sapiens chromosome 1 open reading frame 34 (C1orf34), mRNA
 XM_375732 Homo sapiens similar to Retrovirus-related Pol polyprotein from transposon 1
 XM_375737 Homo sapiens DnaJ (Hsp40) homolog, subfamily C, member 6 (DNAJC6), mRNA
 XM_375738 Homo sapiens hypothetical gene supported by BC047053 (LOC400757), mRNA
 XM_375744 Homo sapiens hypothetical protein LOC339524 (LOC339524), mRNA
 XM_375746 Homo sapiens similar to Interferon-induced guanylate-binding protein 1 (GTF)
 XM_375747 Homo sapiens similar to Interferon-induced guanylate-binding protein 1 (GTF)
 XM_375753 Homo sapiens hypothetical gene supported by AK092728 (LOC400765), mRNA
 XM_375754 Homo sapiens hypothetical protein LOC163404 (LOC163404), mRNA
 XM_375756 Homo sapiens similar to Stromal cell derived factor receptor 2 (LOC400766), mRNA
 XM_375761 Homo sapiens similar to Matrin 3 (LOC400767), mRNA
 XM_375762 Homo sapiens netrin G1 (NTNG1), mRNA
 XM_375770 Homo sapiens leucine-rich repeats and immunoglobulin-like domains 2 (LRI)
 XM_375774 Homo sapiens similar to hypothetical protein DKFZp434A171 (LOC400773), mRNA
 XM_375775 Homo sapiens similar to erbB3 (LOC400774), mRNA
 XM_375779 Homo sapiens similar to hypothetical protein AE2 (LOC400776), mRNA
 XM_375783 Homo sapiens hypothetical protein LOC149013 (LOC149013), mRNA
 XM_375785 Homo sapiens similar to autoantigen La (LOC400779), mRNA
 XM_375802 Homo sapiens hypothetical gene supported by BC014333 (LOC400784), mRNA
 XM_375803 Homo sapiens similar to putative UST1-like organic anion transporter (LOC4)
 XM_375806 Homo sapiens KIAA0476 gene product (KIAA0476), mRNA
 XM_375809 Homo sapiens hypothetical protein LOC126669 (LOC126669), mRNA
 XM_375810 Homo sapiens similar to glucocerebrosidase (LOC400787), mRNA
 XM_375811 Homo sapiens similar to misato (LOC400788), mRNA
 XM_375812 Homo sapiens KIAA0907 protein (KIAA0907), mRNA
 XM_375814 Homo sapiens similar to LINE-1 REVERSE TRANSCRIPTASE HOMOLOG (I
 XM_375816 Homo sapiens similar to Slm17 protein (LOC400792), mRNA
 XM_375821 Homo sapiens hypothetical gene supported by AK128015 (LOC400799), mRNA
 XM_375825 Homo sapiens kinesin family member 14 (KIF14), mRNA
 XM_375833 Homo sapiens hypothetical protein LOC284581 (LOC284581), mRNA
 XM_375834 Homo sapiens inhibitor of kappa light polypeptide gene enhancer in B-cells, I
 XM_375837 Homo sapiens KIAA0205 gene product (KIAA0205), mRNA
 XM_375838 Homo sapiens hypothetical protein LOC128387 (LOC128387), mRNA
 XM_375841 Homo sapiens hypothetical gene supported by AK128488 (LOC400804), mRNA
 XM_375842 Homo sapiens similar to hypothetical protein LOC349114 (LOC400805), mRNA
 XM_375843 Homo sapiens similar to hypothetical protein FLJ25976 (LOC400806), mRNA
 XM_375845 Homo sapiens similar to bA47615.3 (novel protein similar to septin) (LOC40)
 XM_375846 Homo sapiens similar to hypothetical protein FLJ25976 (LOC400808), mRNA
 XM_375848 Homo sapiens KIAA0792 gene product (KIAA0792), mRNA
 XM_375849 Homo sapiens similar to Hypothetical protein CBG08611 (LOC400809), mRNA
 XM_375850 Homo sapiens similar to Ferritin heavy chain (Ferritin H subunit) (LOC40081)
 XM_375851 Homo sapiens KIAA0133 gene product (KIAA0133), mRNA
 XM_375853 Homo sapiens protein BAP28 (FLJ10359), mRNA
 XM_375856 Homo sapiens similar to bA47615.3 (novel protein similar to septin) (LOC40)
 XM_375863 Homo sapiens similar to 60S ribosomal protein L23a (LOC400814), mRNA
 XM_375865 Homo sapiens similar to BC02216 protein (LOC400815), mRNA
 XM_375869 Homo sapiens similar to LOC375080 protein (LOC400818), mRNA
 XM_375873 Homo sapiens similar to KIAA0447 protein (LOC400820), mRNA
 XM_375875 Homo sapiens similar to hypothetical protein LOC349114 (LOC400821), mRNA
 XM_375876 Homo sapiens similar to hypothetical protein FLJ25976 (LOC400822), mRNA
 XM_375882 Homo sapiens similar to chromosome 14 open reading frame 24 (LOC40082)
 XM_375885 Homo sapiens similar to C219-reactive peptide (LOC400824), mRNA
 XM_375887 Homo sapiens similar to Calyculin (Prolactin receptor associated protein) (PF
 XM_375897 Homo sapiens similar to AG1 (LOC400826), mRNA

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XM_375899 Homo sapiens similar to Ba1-651 (LOC400827), mRNA
 XM_375902 Homo sapiens similar to beta-defensin 32 precursor (LOC400830), mRNA
 XM_375904 Homo sapiens similar to RIKEN cDNA 4933425O20 (LOC400833), mRNA
 XM_375911 Homo sapiens KIAA0186 gene product (KIAA0186), mRNA
 XM_375912 Homo sapiens similar to hypothetical protein FLJ38374 (LOC400840), mRNA
 XM_375914 Homo sapiens similar to dJ1184F.4 (novel protein similar to nucleolar prote
 XM_375917 Homo sapiens hypothetical protein LOC149692 (LOC149692), mRNA
 XM_375922 Homo sapiens similar to Protein C20orf85 (LOC400848), mRNA
 XM_375925 Homo sapiens similar to cyclin-like F-box (3A784) (LOC400854), mRNA
 XM_375928 Homo sapiens similar to hypothetical protein MGC30156 (LOC400855), mRNA
 XM_375929 Homo sapiens hypothetical gene supported by AK123815 (LOC400856), mR
 XM_375930 Homo sapiens similar to PRED3 (LOC400857), mRNA
 XM_375931 Homo sapiens similar to PRED4 (LOC400858), mRNA
 XM_375934 Homo sapiens similar to LINE-1 REVERSE TRANSCRIPTASE HOMOLOG (I
 XM_375935 Homo sapiens hypothetical protein LOC284825 (LOC284825), mRNA
 XM_375936 Homo sapiens chaperonin containing TCP1, subunit 8 (theta) (CCT8), mRNA
 XM_375941 Homo sapiens hypothetical gene supported by AK127082 (LOC400867), mR
 XM_375946 Homo sapiens hypothetical gene supported by AK124122 (LOC400878), mR
 XM_375948 Homo sapiens similar to Serine/threonine-protein kinase Nek2 (NIMA-related
 XM_375951 Homo sapiens similar to Gene with similarity to rat kidney-specific (KS) gene
 XM_375953 Homo sapiens hypothetical gene supported by AK129567; NM_201401 (LOC
 XM_375954 Homo sapiens similar to LOC284861 protein (LOC400887), mRNA
 XM_375955 Homo sapiens similar to immunoglobulin superfamily, member 3; immunoglo
 XM_375958 Homo sapiens similar to proline dehydrogenase (oxidase) 1; proline oxidase
 XM_375963 Homo sapiens similar to hypothetical protein LOC145497 (LOC400891), mR
 XM_375964 Homo sapiens similar to breakpoint cluster region isoform 1 (LOC400892), r
 XM_375965 Homo sapiens similar to Gamma-glutamyltranspeptidase 1 precursor (Gamm
 XM_375966 Homo sapiens similar to immunoglobulin superfamily, member 3; immunoglo
 XM_375993 Homo sapiens similar to a putative protein with homology to a sequence betw
 XM_375996 Homo sapiens similar to Gamma-glutamyltransferase-like protein 4 (LOC40
 XM_375997 Homo sapiens similar to breakpoint cluster region isoform 1 (LOC400918), r
 XM_376001 Homo sapiens similar to low density lipoprotein receptor-related protein 5; lo
 XM_376003 Homo sapiens hypothetical gene supported by AK056895 (LOC400924), mR
 XM_376007 Homo sapiens KIAA0645 gene product (KIAA0645), mRNA
 XM_376008 Homo sapiens hypothetical protein LOC91464 (LOC91464), mRNA
 XM_376010 Homo sapiens similar to TPTE and PTEN homologous inositol lipid phosphat
 XM_376013 Homo sapiens hypothetical protein LOC200321 (LOC200321), mRNA
 XM_376018 Homo sapiens KIAA1644 protein (KIAA1644), mRNA
 XM_376019 Homo sapiens similar to hypothetical protein (LOC400930), mRNA
 XM_376020 Homo sapiens hypothetical gene supported by AK130875 (LOC400931), mR
 XM_376021 Homo sapiens hypothetical gene supported by AK128136 (LOC400932), mR
 XM_376022 Homo sapiens hypothetical gene supported by AK126356 (LOC400934), mR
 XM_376023 Homo sapiens zinc finger, BED domain containing 4 (ZBED4), mRNA
 XM_376024 Homo sapiens similar to interleukin 17 receptor E; EST AA589509 (LOC400
 XM_376031 Homo sapiens hypothetical gene supported by AK123041 (LOC400940), mR
 XM_376032 Homo sapiens hypothetical gene supported by AK124409 (LOC400941), mR
 XM_376033 Homo sapiens hypothetical protein LOC339789 (LOC339789), mRNA
 XM_376034 Homo sapiens similar to AILT5830 (LOC400943), mRNA
 XM_376043 Homo sapiens similar to RIKEN cDNA 2310016E02 (LOC400948), mRNA
 XM_376044 Homo sapiens SPTF-associated factor 65 gamma (STAF65(gamma)), mRNA/
 XM_376048 Homo sapiens similar to FKSG60 (LOC400949), mRNA
 XM_376049 Homo sapiens hypothetical gene supported by AK124893 (LOC400950), mR
 XM_376051 Homo sapiens similar to NGNL6975 (LOC400952), mRNA
 XM_376056 Homo sapiens similar to echinoderm microtubule associated protein like 5 (L
 XM_376059 Homo sapiens SERTA domain containing 2 (SERTAD2), mRNA
 XM_376060 Homo sapiens KIAA0053 gene product (KIAA0053), mRNA
 XM_376062 Homo sapiens similar to KIAA1155 protein (LOC400961), mRNA

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XM_376068 Homo sapiens similar to LP3727 (LOC400962), mRNA
 XM_376072 Homo sapiens hypothetical gene supported by AK098018 (LOC400965), mR
 XM_376073 Homo sapiens similar to Ig kappa variable region (LOC400967), mRNA
 XM_376074 Homo sapiens LOC400968 (LOC400968), mRNA
 XM_376094 Homo sapiens hypothetical protein LOC90499 (LOC90499), mRNA
 XM_376097 Homo sapiens similar to seven transmembrane helix receptor (LOC400964),
 XM_376099 Homo sapiens similar to Glycerol-3-phosphate acyltransferase, mitochondria
 XM_376100 Homo sapiens similar to KIAA1641 protein (LOC400986), mRNA
 XM_376101 Homo sapiens similar to LOC375251 protein (LOC400987), mRNA
 XM_376106 Homo sapiens similar to RIKEN cDNA 6330578E17 (LOC400989), mRNA
 XM_376108 Homo sapiens similar to Sodium/hydrogen exchanger 4 (Na⁺/H⁺ exchang
 XM_376111 Homo sapiens plasminogen-related protein A (LOC285189), mRNA
 XM_376112 Homo sapiens similar to RAN-binding protein 2-like 1 isoform 1; sperm meml
 XM_376117 Homo sapiens hypothetical gene supported by AK095987 (LOC400994), mR
 XM_376118 Homo sapiens hypothetical gene supported by AK095987 (LOC400996), mR
 XM_376121 Homo sapiens hypothetical gene supported by AK095987 (LOC400998), mR
 XM_376125 Homo sapiens similar to Single-stranded DNA-binding protein, isoform b (LO
 XM_376126 Homo sapiens similar to hypothetical protein A230046P18 (LOC401003), mF
 XM_376127 Homo sapiens similar to pole protein; Expressed in prostate, ovary, testis, ar
 XM_376130 Homo sapiens similar to pole protein; Expressed in prostate, ovary, testis, ar
 XM_376139 Homo sapiens hypothetical gene supported by AK123815 (LOC401011), mR
 XM_376141 Homo sapiens similar to zinc finger protein 285 (LOC401012), mRNA
 XM_376142 Homo sapiens hypothetical gene supported by AK057980; AK092189 (LOC4
 XM_376144 Homo sapiens hypothetical protein LOC339745 (LOC339745), mRNA
 XM_376148 Homo sapiens similar to RIKEN cDNA 5830415L20 (LOC401015), mRNA
 XM_376150 Homo sapiens similar to ribosomal protein S3a; 40S ribosomal protein S3a; v
 XM_376154 Homo sapiens similar to ribosomal protein S15; rat insulinoma gene (LOC40
 XM_376158 Homo sapiens hypothetical gene supported by AK126104; BX648733 (LOC4
 XM_376160 Homo sapiens similar to CDNA sequence BC030440 (LOC401026), mRNA
 XM_376161 Homo sapiens similar to IIDS6411 (LOC401027), mRNA
 XM_376182 Homo sapiens similar to Nucleophosmin 1 (LOC401028), mRNA
 XM_376165 Homo sapiens similar to DAZ associated protein 2; deleted in azoospermia a
 XM_376171 Homo sapiens KIAA1843 protein (KIAA1843), mRNA
 XM_376172 Homo sapiens zinc finger protein 142 (clone pHZ-49) (ZNF142), mRNA
 XM_376178 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA
 XM_376179 Homo sapiens LOC401034 (LOC401034), mRNA
 XM_376180 Homo sapiens similar to archease (LOC401035), mRNA
 XM_376185 Homo sapiens similar to ankyrin repeat and SOCS box-containing 18; SOCS
 XM_376188 Homo sapiens hypothetical protein LOC93463 (LOC93463), mRNA
 XM_376189 Homo sapiens DKFZP586K1520 protein (DKFZP586K1520), mRNA
 XM_376190 Homo sapiens hypothetical gene supported by AK125867 (LOC401039), mR
 XM_376191 Homo sapiens hypothetical gene supported by AK127861 (LOC401040), mR
 XM_376193 Homo sapiens FERM, RhoGEF and pleckstrin domain protein 2 (FARF2), ml
 XM_376195 Homo sapiens hypothetical gene supported by AK123321 (LOC401045), mR
 XM_376200 Homo sapiens similar to inhibitor of growth family, member 5 (LOC401047), i
 XM_376201 Homo sapiens ER degradation enhancer, mannosidase alpha-like 1 (EDEM1
 XM_376203 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA
 XM_376206 Homo sapiens hypothetical protein LOC285375 (LOC285375), mRNA
 XM_376207 Homo sapiens similar to FLJ00274 protein (LOC401054), mRNA
 XM_376209 Homo sapiens similar to Nonhistone chromosomal protein HMG-17 (High-mc
 XM_376212 Homo sapiens hypothetical protein LOC339862 (LOC339862), mRNA
 XM_376225 Homo sapiens similar to serine protease-like 1 (LOC401053), mRNA
 XM_376227 Homo sapiens hypothetical gene supported by AK097724 (LOC401064), mR
 XM_376232 Homo sapiens Vpr-binding protein (VprBP), mRNA
 XM_376233 Homo sapiens similar to hypothetical protein MGC39725 (LOC401067), mR
 XM_376238 Homo sapiens hypothetical protein LOC285331 (LOC285331), mRNA
 XM_376239 Homo sapiens similar to FtsJ homolog 2 isoform b; cell division protein FtsJ;

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XM_376241 Homo sapiens hypothetical gene supported by AK125779 (LOC401070), mR
 XM_376243 Homo sapiens hypothetical gene supported by AK125942 (LOC401072), mR
 XM_376247 Homo sapiens similar to double homeobox protein (LOC401074), mRNA
 XM_376248 Homo sapiens similar to zinc finger protein 39 (LOC401075), mRNA
 XM_376249 Homo sapiens similar to CAP, adenylate cyclase-associated protein 1; aden
 XM_376254 Homo sapiens hypothetical protein DKFZp667G2110 (DKFZp667G2110), m
 XM_376256 Homo sapiens hypothetical gene supported by AK126064 (LOC401080), m
 XM_376257 Homo sapiens similar to hypothetical protein FLJ25976 (LOC401082), mR/
 XM_376258 Homo sapiens similar to Nuclear transcription factor Y subunit beta (NF-Y pr
 XM_376267 Homo sapiens similar to hypothetical protein, MNCb-4779 (LOC401087), mR
 XM_376268 Homo sapiens hypothetical gene supported by AK127796 (LOC401088), mR
 XM_376269 Homo sapiens hypothetical gene supported by AK125319 (LOC401089), mR
 XM_376278 Homo sapiens similar to RIKEN cDNA 0610027B03 (LOC401095), mRNA
 XM_376280 Homo sapiens hypothetical protein BC010062 (LOC152078), mRNA
 XM_376281 Homo sapiens hypothetical gene supported by BC031660 (LOC401097), mR
 XM_376284 Homo sapiens hypothetical protein BC011266 (LOC935566), mRNA
 XM_376287 Homo sapiens hypothetical gene supported by AK128090 (LOC401100), mR
 XM_376290 Homo sapiens hypothetical gene supported by AK124384 (LOC401105), mR
 XM_376292 Homo sapiens hypothetical gene supported by AK129507 (LOC401109), mR
 XM_376299 Homo sapiens hypothetical gene supported by AK093135 (LOC401114), mR
 XM_376300 Homo sapiens hypothetical gene supported by AK124538 (LOC401116), mR
 XM_376301 Homo sapiens similar to chromosome 11 open reading frame2; chromosome
 XM_376303 Homo sapiens hypothetical protein LOC285484 (LOC285484), mRNA
 XM_376305 Homo sapiens similar to double ubiquitinating enzyme 3 (LOC401121), mRNA
 XM_376306 Homo sapiens similar to hypothetical protein LOC169270 (LOC401122), mR/
 XM_376307 Homo sapiens hypothetical protein DKFZp667E0512 (DKFZp667E0512), mF
 XM_376309 Homo sapiens hypothetical protein LOC285540 (LOC285540), mRNA
 XM_376310 Homo sapiens zinc finger, CCHC domain containing 4 (ZCCHC4), mRNA
 XM_376312 Homo sapiens hypothetical gene supported by AK127623 (LOC401123), mR
 XM_376314 Homo sapiens similar to TBC1 domain family member 1 (LOC401125), mR/
 XM_376317 Homo sapiens similar to SNAG1 (LOC401130), mRNA
 XM_376318 Homo sapiens similar to hypothetical protein FLJ30672 (LOC401132), mR/
 XM_376320 Homo sapiens similar to RIKEN cDNA 9930019B18 gene (LOC401136), mR/
 XM_376322 Homo sapiens similar to hypothetical protein (LOC401137), mRNA
 XM_376323 Homo sapiens similar to RST1689 (LOC401138), mRNA
 XM_376324 Homo sapiens similar to Ameloblastin precursor (LOC401139), mRNA
 XM_376325 Homo sapiens hypothetical protein FLJ13105 (FLJ13105), mRNA
 XM_376327 Homo sapiens similar to hypothetical protein LOC231503 (LOC401141), mR/
 XM_376328 Homo sapiens family with sequence similarity 13, member A1 (FAM13A1), m
 XM_376331 Homo sapiens KIAA1680 protein (KIAA1680), mRNA
 XM_376333 Homo sapiens similar to elongation factor 1 alpha (LOC401146), mRNA
 XM_376334 Homo sapiens similar to hypothetical protein (LOC401147), mRNA
 XM_376338 Homo sapiens hypothetical gene supported by AK127273 (LOC401150), mR
 XM_376339 Homo sapiens similar to RIKEN cDNA 1810037117 (LOC401152), mRNA
 XM_376342 Homo sapiens similar to bA291L22.2 (similar to CDC10 (cell division cycle 1
 XM_376347 Homo sapiens hypothetical gene supported by AK126441 (LOC401157), mR
 XM_376348 Homo sapiens similar to hypothetical protein (L1H 3 region) - human (LOC4
 XM_376349 Homo sapiens hypothetical protein LOC201725 (LOC201725), mRNA
 XM_376350 Homo sapiens PDZ domain containing guanine nucleotide exchange factor (i
 XM_376353 Homo sapiens similar to hypothetical protein FLJ20035 (LOC401160), mR/
 XM_376354 Homo sapiens similar to inactive progesterone receptor, 23 kD; likely ortho
 XM_376355 Homo sapiens KIAA0626 gene product (KIAA0626), mRNA
 XM_376364 Homo sapiens hypothetical gene supported by AK126844 (LOC401166), mR
 XM_376365 Homo sapiens hypothetical protein BC014011 (LOC116349), mRNA
 XM_376366 Homo sapiens DKFZP564I1171 protein (DKFZP564I1171), mRNA
 XM_376368 Homo sapiens similar to Programmed cell death protein 6 (Probable calcium
 XM_376370 Homo sapiens hypothetical gene supported by AK090679 (LOC401172), mR

XM_376371 Homo sapiens similar to hypothetical protein FLJ36144 (LOC401174), mRNA
 XM_376372 Homo sapiens hypothetical protein LOC134121 (LOC134121), mRNA
 XM_376376 Homo sapiens similar to SMA3 protein (LOC401179), mRNA
 XM_376379 Homo sapiens similar to RIKEN cDNA 4921505C17 (LOC401183), mRNA
 XM_376383 Homo sapiens similar to cAMP-specific phosphodiesterase PDE407 (LOC40
 XM_376386 Homo sapiens similar to DNA segment, Chr 13, Brigham & Womens Genetic
 XM_376387 Homo sapiens hypothetical gene supported by AK127903 (LOC401191), mR
 XM_376389 Homo sapiens similar to small EDRK-rich factor 1A, telomeric; spinal muscu
 XM_376391 Homo sapiens similar to psi neuronal apoptosis inhibitory protein (LOC4011
 XM_376394 Homo sapiens hypothetical gene supported by AK130705 (LOC401195), mR
 XM_376395 Homo sapiens similar to POM121 membrane glycoprotein-like 1 (LOC401191
 XM_376397 Homo sapiens hypothetical protein LOC153561 (LOC153561), mRNA
 XM_376403 Homo sapiens similar to ribosomal protein L7-like 1 (LOC401197), mRNA
 XM_376405 Homo sapiens Rho-guanine nucleotide exchange factor (RGNEF), mRNA
 XM_376412 Homo sapiens similar to KIAA0825 protein (LOC401202), mRNA
 XM_376413 Homo sapiens hypothetical protein DKFZp564C0469 (DKFZp564C0469), mR
 XM_376416 Homo sapiens similar to Beta-glucuronidase precursor (Beta-G1) (LOC4012
 XM_376419 Homo sapiens hypothetical protein LOC285638 (LOC285638), mRNA
 XM_376420 Homo sapiens similar to 40S ribosomal protein S25 (LOC401206), mRNA
 XM_376423 Homo sapiens hypothetical gene supported by AK126569 (LOC401207), mR
 XM_376427 Homo sapiens LOC401208 (LOC401208), mRNA
 XM_376428 Homo sapiens similar to HYPOTHETICAL PROTEIN ORF-1137 (LOC40120
 XM_376430 Homo sapiens similar to nuclear receptor coactivator 4; RET-activating gene
 XM_376433 Homo sapiens hypothetical protein LOC153218 (LOC153218), mRNA
 XM_376436 Homo sapiens hypothetical protein LOC134466 (LOC134466), mRNA
 XM_376440 Homo sapiens hypothetical protein LOC285629 (LOC285629), mRNA
 XM_376443 Homo sapiens hypothetical gene supported by AK097772 (LOC401217), mR
 XM_376444 Homo sapiens hypothetical protein LOC133491 (LOC133491), mRNA
 XM_376447 Homo sapiens similar to hypothetical protein (LOC401221), mRNA
 XM_376453 Homo sapiens similar to KIAA0752 protein (LOC401223), mRNA
 XM_376454 Homo sapiens similar to acetoacetyl-CoA synthetase; acetoacetate-CoA liga
 XM_376458 Homo sapiens hypothetical gene supported by AK093729; AK128780; BX64'
 XM_376461 Homo sapiens similar to bA476115.3 (novel protein similar to septin) (LOC40
 XM_376463 Homo sapiens hypothetical protein MGC39372 (MGC39372), mRNA
 XM_376464 Homo sapiens similar to HIV TAT specific factor 1; cofactor required for Tat s
 XM_376469 Homo sapiens hypothetical gene supported by AK026805 (LOC401236), mR
 XM_376471 Homo sapiens similar to chromosome 15 open reading frame 2 (LOC401238
 XM_376472 Homo sapiens similar to KIAA0319 (LOC401239), mRNA
 XM_376473 Homo sapiens similar to SMA3-like protein bA239L20.1 (LOC401240), mRNA
 XM_376474 Homo sapiens Integral membrane glycoprotein-like (LOC166994), mRNA
 XM_376479 Homo sapiens mediator of DNA damage checkpoint 1 (MDC1), mRNA
 XM_376480 Homo sapiens hypothetical gene supported by AK098012 (LOC401247), mR
 XM_376486 Homo sapiens similar to coiled-coil domain 1 protein precursor (LOC401250;
 XM_376487 Homo sapiens similar to NG23 (LOC401251), mRNA
 XM_376488 Homo sapiens hypothetical gene supported by AK123889 (LOC401252), mR
 XM_376491 Homo sapiens hypothetical gene supported by AK125740 (LOC401253), mR
 XM_376498 Homo sapiens similar to kinesin-related protein 3A (LOC401259), mRNA
 XM_376499 Homo sapiens hypothetical gene supported by AK123643 (LOC401260), mR
 XM_376503 Homo sapiens ectonucleotide pyrophosphatase/phosphodiesterase 4 (putati
 XM_376505 Homo sapiens similar to hypothetical protein FLJ30295 (LOC401263), mRNA
 XM_376508 Homo sapiens hypothetical gene supported by AK091177 (LOC401265), mR
 XM_376516 Homo sapiens myosin VI (MYO6), mRNA
 XM_376518 Homo sapiens chromosome 6 open reading frame 84 (C8orf84), mRNA
 XM_376519 Homo sapiens ankyrin repeat domain 6 (ANKRD6), mRNA
 XM_376522 Homo sapiens similar to Heat shock protein 67B2 (LOC401270), mRNA
 XM_376525 Homo sapiens zinc finger protein 450 (ZNF450), mRNA
 XM_376527 Homo sapiens hypothetical gene supported by AK124171 (LOC401271), mR

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XM_376532 Homo sapiens similar to KIAA0408 protein (LOC401272), mRNA
 XM_376533 Homo sapiens similar to FLJ4670 protein (LOC401273), mRNA
 XM_376535 Homo sapiens chromosome 6 open reading frame 207 (C6orf207), mRNA
 XM_376536 Homo sapiens similar to hypothetical protein (LOC401274), mRNA
 XM_376537 Homo sapiens BCL2-associated transcription factor 1 (BCLAF1), mRNA
 XM_376540 Homo sapiens chromosome 6 open reading frame 56 (C6orf56), mRNA
 XM_376541 Homo sapiens hypothetical gene supported by AK126903 (LOC401278), mRNA
 XM_376547 Homo sapiens RNA binding motif protein 16 (RBM16), mRNA
 XM_376549 Homo sapiens hypothetical gene supported by AK125637 (LOC401280), mRNA
 XM_376550 Homo sapiens KIAA1423 (KIAA1423), mRNA
 XM_376554 Homo sapiens similar to T-complex protein 10A homolog (LOC401285), mRNA
 XM_376555 Homo sapiens hypothetical gene supported by AK127120 (LOC401286), mRNA
 XM_376556 Homo sapiens chromosome 6 open reading frame 70 (C6orf70), mRNA
 XM_376557 Homo sapiens hypothetical gene supported by AK056013 (LOC401288), mRNA
 XM_376558 Homo sapiens hypothetical gene supported by AK127120 (LOC401293), mRNA
 XM_376560 Homo sapiens hypothetical gene supported by AK125637 (LOC401295), mRNA
 XM_376564 Homo sapiens unc-84 homolog A (C. elegans) (UNC84A), mRNA
 XM_376565 Homo sapiens hypothetical gene supported by BC031661 (LOC401298), mRNA
 XM_376566 Homo sapiens hypothetical protein LOC285924 (LOC285924), mRNA
 XM_376567 Homo sapiens KIAA1856 protein (KIAA1856), mRNA
 XM_376568 Homo sapiens hypothetical gene supported by AK125308 (LOC401300), mRNA
 XM_376569 Homo sapiens hypothetical gene supported by AK125353 (LOC401302), mRNA
 XM_376571 Homo sapiens ubiquitin specific protease 42 (USP42), mRNA
 XM_376573 Homo sapiens similar to beta-1,4-mannosyltransferase; beta-1,4 mannosyltr
 XM_376575 Homo sapiens hypothetical gene supported by AK027125 (LOC401307), mRNA
 XM_376576 Homo sapiens similar to Chain , Heat-Shock Cognate 70kd Protein (44kd Atq
 XM_376577 Homo sapiens similar to heat shock 70kDa protein 8 isoform 2; heat shock c
 XM_376578 Homo sapiens PHD finger protein 14 (PHF14), mRNA
 XM_376585 Homo sapiens similar to Dual specificity protein kinase CLK2 (CDC like kinas
 XM_376586 Homo sapiens similar to Hypothetical protein KIAA0087 (HA1002) (LOC4013
 XM_376587 Homo sapiens similar to mKIAA0038 protein (LOC401316), mRNA
 XM_376588 Homo sapiens KIAA0644 gene product (KIAA0644), mRNA
 XM_376589 Homo sapiens KIAA0241 protein (KIAA0241), mRNA
 XM_376590 Homo sapiens LOC89231 (LOC89231), mRNA
 XM_376591 Homo sapiens similar to KIAA0877 protein (LOC401322), mRNA
 XM_376593 Homo sapiens similar to RIKEN cDNA 9330128H10 gene (LOC401323), mRNA
 XM_376595 Homo sapiens hypothetical gene supported by AF447883 (LOC401325), mRNA
 XM_376597 Homo sapiens similar to sequence-specific single-stranded-DNA-binding pro
 XM_376598 Homo sapiens similar to t-complex 1; T-complex locus TCP-1; t-complex 1 (a
 XM_376600 Homo sapiens similar to RAS p21 protein activator 4; GTPase activating prot
 XM_376602 Homo sapiens similar to Ras GTPase-activating protein 4 (RasGAP-activatin
 XM_376604 Homo sapiens similar to cell division cycle 10 homolog (LOC401332), mRNA
 XM_376605 Homo sapiens similar to hypothetical protein FLJ25976 (LOC401333), mRNA
 XM_376607 Homo sapiens hypothetical gene supported by AK126096 (LOC401335), mRNA
 XM_376609 Homo sapiens growth factor receptor-bound protein 10 (GRB10), mRNA
 XM_376610 Homo sapiens hypothetical gene supported by AK097404; NM_198284 (LOC
 XM_376611 Homo sapiens hypothetical gene supported by AK127870 (LOC401337), mRNA
 XM_376612 Homo sapiens similar to SMT3 suppressor of mif two 3 homolog 2 (LOC4013
 XM_376613 Homo sapiens similar to hypothetical protein DKFp434F142 (LOC401341), mRNA
 XM_376614 Homo sapiens LOC401350 (LOC401350), mRNA
 XM_376615 Homo sapiens hypothetical gene supported by BC040831 (LOC401351), mRNA
 XM_376616 Homo sapiens hypothetical gene supported by BC040831 (LOC401354), mRNA
 XM_376617 Homo sapiens similar to BC060615 protein (LOC401355), mRNA
 XM_376618 Homo sapiens similar to CAGL79 (LOC401356), mRNA
 XM_376619 Homo sapiens similar to hypothetical protein LOC285908 (LOC401357), mRNA
 XM_376621 Homo sapiens similar to hypothetical protein FLJ25037 (LOC401360), mRNA
 XM_376622 Homo sapiens similar to hypothetical protein MGC16733 similar to CG12113

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XM_376623 Homo sapiens similar to MGC16733 protein (LOC401362), mRNA
 XM_376625 Homo sapiens similar to hypothetical protein FLJ10900 (LOC401369), mRNA
 XM_376626 Homo sapiens similar to Williams Beuren syndrome chromosome region 19 i
 XM_376628 Homo sapiens similar to Neutrophil cytosol factor 1 (NCF-1) (Neutrophil NAD
 XM_376629 Homo sapiens similar to transcription factor GTF2IRD2 (LOC401375), mRNA
 XM_376630 Homo sapiens similar to Nuclear envelope pore membrane protein POM 121
 XM_376631 Homo sapiens Williams Beuren syndrome chromosome region 24 (WBSCR2
 XM_376636 Homo sapiens similar to PMS4 (LOC401378), mRNA
 XM_376638 Homo sapiens similar to PMS4 (LOC401379), mRNA
 XM_376639 Homo sapiens similar to Williams Beuren syndrome chromosome region 19 i
 XM_376640 Homo sapiens similar to PMS4 homolog mismatch repair protein - human (Li
 XM_376642 Homo sapiens tripartite motif-containing 50B (TRIM50B), mRNA
 XM_376643 Homo sapiens similar to hypothetical protein LOC285908 (LOC401383), mRi
 XM_376647 Homo sapiens sema domain, immunoglobulin domain (Ig), short basic doma
 XM_376648 Homo sapiens similar to hypothetical protein 4932412H11 (LOC401387), mF
 XM_376649 Homo sapiens hypothetical protein FLJ39885 (FLJ39885), mRNA
 XM_376651 Homo sapiens hypothetical gene supported by AK124274 (LOC401388), mR
 XM_376652 Homo sapiens distal-less homeo box 6 (DLX6), mRNA
 XM_376653 Homo sapiens similar to chromosome 11 open reading frame2; chromosome
 XM_376655 Homo sapiens similar to importin alpha 1b (LOC401391), mRNA
 XM_376656 Homo sapiens hypothetical protein FLJ22037 (FLJ22037), mRNA
 XM_376657 Homo sapiens hypothetical protein LOC285889 (LOC285889), mRNA
 XM_376658 Homo sapiens similar to Zinc-alpha-2-glycoprotein precursor (Zn-alpha-2-gly
 XM_376663 Homo sapiens similar to reverse transcriptase related protein (LOC401395),
 XM_376664 Homo sapiens KIAA1218 protein (KIAA1218), mRNA
 XM_376665 Homo sapiens hypothetical protein LOC286009 (LOC286009), mRNA
 XM_376668 Homo sapiens similar to Serine/threonine-protein kinase tousled-like 2 (Tous
 XM_376670 Homo sapiens similar to hypothetical protein FLJ25976 (LOC401402), mRNA
 XM_376671 Homo sapiens coatomer protein complex, subunit gamma 2 (COPG2), mRN
 XM_376672 Homo sapiens similar to ribosomal protein S14 (LOC401404), mRNA
 XM_376677 Homo sapiens hypothetical protein LOC155006 (LOC155006), mRNA
 XM_376679 Homo sapiens hypothetical protein FLJ25778 (FLJ25778), mRNA
 XM_376680 Homo sapiens KIAA1718 protein (KIAA1718), mRNA
 XM_376681 Homo sapiens similar to RAB19, member RAS oncogene family (LOC401401
 XM_376683 Homo sapiens LCHN protein (LCHN), mRNA
 XM_376684 Homo sapiens hypothetical protein LOC93432 (LOC93432), mRNA
 XM_376707 Homo sapiens similar to KIAA0738 protein (LOC401426), mRNA
 XM_376712 Homo sapiens FLJ43692 protein (FLJ43692), mRNA
 XM_376713 Homo sapiens similar to Olfactory receptor 2A7 (LOC401427), mRNA
 XM_376715 Homo sapiens similar to seven transmembrane helix receptor (LOC401428),
 XM_376716 Homo sapiens similar to KIAA1285 protein (LOC401429), mRNA
 XM_376717 Homo sapiens likely ortholog of mouse zinc finger protein EZI (EZI), mRNA
 XM_376718 Homo sapiens FLJ45737 protein (FLJ45737), mRNA
 XM_376719 Homo sapiens similar to KIAA2036 protein (LOC401430), mRNA
 XM_376720 Homo sapiens KIAA0543 protein (KIAA0543), mRNA
 XM_376722 Homo sapiens hypothetical protein LOC155036 (LOC155036), mRNA
 XM_376724 Homo sapiens KIAA1402 protein (CSG1cA-T), mRNA
 XM_376725 Homo sapiens hypothetical gene supported by AK127717 (LOC401433), mR
 XM_376727 Homo sapiens hypothetical protein LOC285888 (LOC285888), mRNA
 XM_376728 Homo sapiens hypothetical protein LOC155435 (LOC155435), mRNA
 XM_376730 Homo sapiens ubiquitin-protein isopeptide ligase (E3) (KIAA0010), mRNA
 XM_376736 Homo sapiens similar to Hypothetical protein KIAA0711 (LOC401444), mRN
 XM_376741 Homo sapiens similar to ubiquitin-specific protease 17-like protein (LOC4014
 XM_376746 Homo sapiens similar to seven transmembrane helix receptor (LOC401450),
 XM_376750 Homo sapiens similar to chromosome 11 open reading frame2; chromosome
 XM_376754 Homo sapiens similar to deubiquitinating enzyme 3 (LOC401453), mRNA
 XM_376756 Homo sapiens similar to hypothetical protein SB153 isoform 1 (LOC401454),

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XM_376757 Homo sapiens similar to KIAA1456 protein (LOC401455), mRNA
 XM_376761 Homo sapiens hypothetical gene supported by BC062364; BX047289 (LOC4
 XM_376763 Homo sapiens similar to TRANSCRIPTION FACTOR COE2 (EARLY B-CELL
 XM_376764 Homo sapiens paraneoplastic antigen MA2 (PNMA2), mRNA
 XM_376771 Homo sapiens hypothetical gene supported by AK128232 (LOC401459), mR
 XM_376774 Homo sapiens similar to hypothetical protein 4932417K07 (LOC401460), mR
 XM_376776 Homo sapiens thymus high mobility group box protein TOX (TOX), mRNA
 XM_376780 Homo sapiens similar to Myelin P2 protein (LOC401465), mRNA
 XM_376781 Homo sapiens solute carrier family 10 (sodium/bile acid cotransporter family)
 XM_376783 Homo sapiens hypothetical gene supported by BC055062 (LOC401466), mR
 XM_376784 Homo sapiens similar to prot GOR (LOC401467), mRNA
 XM_376785 Homo sapiens similar to NFS1 nitrogen fixation 1 isoform b precursor; cystei
 XM_376786 Homo sapiens similar to Ceruloplasmin precursor (Ferroxidase) (LOC401468
 XM_376787 Homo sapiens similar to 40S ribosomal protein S26 (LOC401470), mRNA
 XM_376791 Homo sapiens hypothetical gene supported by AK127183 (LOC401472), mR
 XM_376793 Homo sapiens similar to RIKEN cDNA A830094I09 gene (LOC401474), mR
 XM_376795 Homo sapiens hypothetical gene supported by AK127771 (LOC401478), mR
 XM_376797 Homo sapiens similar to KIAA0870 protein (LOC401479), mRNA
 XM_376800 Homo sapiens similar to Hypothetical zinc finger protein KIAA0628 (LOC401-
 XM_376801 Homo sapiens hypothetical gene supported by AK091211; AK125852 (LOC4
 XM_376809 Homo sapiens similar to ba110H4.2 (similar to membrane protein) (LOC401-
 XM_376810 Homo sapiens similar to dJ28124.1.2 (Spinal Muscular Atrophy region) (SMA3
 XM_376814 Homo sapiens similar to DDX11 protein (LOC401487), mRNA
 XM_376819 Homo sapiens similar to RIKEN cDNA 4933428I03 (LOC401494), mRNA
 XM_376821 Homo sapiens chromosome 9 open reading frame 14 (C9orf14), mRNA
 XM_376822 Homo sapiens similar to PRO2738 (LOC401497), mRNA
 XM_376824 Homo sapiens similar to RIKEN cDNA A930001M12 gene (LOC401498), mR
 XM_376829 Homo sapiens hypothetical protein LOC158381 (LOC158381), mRNA
 XM_376830 Homo sapiens KIAA0258 (KIAA0258), mRNA
 XM_376833 Homo sapiens LOC401505 (LOC401505), mRNA
 XM_376838 Homo sapiens hypothetical gene supported by AK127145 (LOC401507), mR
 XM_376840 Homo sapiens hypothetical gene supported by AK127145 (LOC401508), mR
 XM_376841 Homo sapiens similar to DKFZP572C163 protein (LOC401509), mRNA
 XM_376843 Homo sapiens similar to LOC286286 protein (LOC401510), mRNA
 XM_376846 Homo sapiens similar to DKFZP572C163 protein (LOC401511), mRNA
 XM_376847 Homo sapiens hypothetical gene supported by AK124538 (LOC401512), mR
 XM_376848 Homo sapiens hypothetical gene supported by AK124538 (LOC401514), mR
 XM_376850 Homo sapiens hypothetical gene supported by AK124122 (LOC401515), mR
 XM_376852 Homo sapiens similar to ba251O17.3 (similar to aquaporin 7) (LOC401516)
 XM_376855 Homo sapiens similar to keratinocyte growth factor-like protein, group III - hu
 XM_376858 Homo sapiens similar to breast cancer antigen NY-BR-1 (LOC401519), mR
 XM_376861 Homo sapiens LOC401520 (LOC401520), mRNA
 XM_376863 Homo sapiens similar to Keratinocyte growth factor precursor (KGF) (Fibrobl
 XM_376866 Homo sapiens similar to ba251O17.3 (similar to aquaporin 7) (LOC401524)
 XM_376869 Homo sapiens similar to tumor suppressor deleted in oral cancer-related 1 (L
 XM_376872 Homo sapiens LOC401529 (LOC401529), mRNA
 XM_376874 Homo sapiens chromosome 9 open reading frame 71 (C9orf71), mRNA
 XM_376876 Homo sapiens similar to coiled-coil-helix-coiled-coil-helix domain containing ;
 XM_376880 Homo sapiens hypothetical gene supported by AK127390 (LOC401534), mR
 XM_376885 Homo sapiens hypothetical gene supported by AK127445 (LOC401535), mR
 XM_376888 Homo sapiens similar to Laminin receptor 1 (LOC401537), mRNA
 XM_376890 Homo sapiens similar to CG14980-PB (LOC375601), mRNA
 XM_376892 Homo sapiens similar to hypothetical protein (LOC401541), mRNA
 XM_376895 Homo sapiens zinc finger protein 510 (ZNF510), mRNA
 XM_376898 Homo sapiens thioredoxin domain containing 4 (endoplasmic reticulum) (TX)
 XM_376899 Homo sapiens similar to RIKEN cDNA D630039A03 gene (LOC401546), mR
 XM_376900 Homo sapiens similar to KIAA0563 gene product (LOC401547), mRNA

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XM_376902 Homo sapiens similar to RIKEN cDNA 4732481H14 (LOC401548), mRNA
 XM_376903 Homo sapiens KIAA0674 protein (KIAA0674), mRNA
 XM_376905 Homo sapiens EGF-like-domain, multiple 5 (EGFL5), mRNA
 XM_376909 Homo sapiens similar to hypothetical protein FLJ25955 (LOC401551), mRNA
 XM_376917 Homo sapiens far upstream element (FUSE) binding protein 3 (FUBP3), mRNA
 XM_376921 Homo sapiens hypothetical protein MGC43306 (MGC43306), mRNA
 XM_376924 Homo sapiens chromosome 9 open reading frame 62 (C9orf62), mRNA
 XM_376925 Homo sapiens similar to DNL zinc finger (3D41) (LOC401560), mRNA
 XM_376930 Homo sapiens hypothetical gene supported by AK127160 (LOC401562), mRNA
 XM_376931 Homo sapiens hypothetical gene supported by AK093587; AK124899 (LOC401563), mRNA
 XM_376939 Homo sapiens similar to 4931415M17 protein (LOC401565), mRNA
 XM_376947 Homo sapiens similar to surfeit 5 isoform b; surfeit locus protein 5 (LOC401566), mRNA
 XM_376949 Homo sapiens similar to Surfeit locus protein 4 (LOC401567), mRNA
 XM_376950 Homo sapiens similar to MGC43306 protein (LOC401568), mRNA
 XM_376960 Homo sapiens similar to RIKEN cDNA 2900002H16 (LOC401569), mRNA
 XM_376965 Homo sapiens LOC401570 (LOC401570), mRNA
 XM_376968 Homo sapiens hypothetical gene supported by AK124122 (LOC401572), mRNA
 XM_376978 Homo sapiens similar to hypothetical protein FLJ33610 (LOC401576), mRNA
 XM_376981 Homo sapiens similar to hypothetical protein (L1H 3 region) - human (LOC401577), mRNA
 XM_376986 Homo sapiens similar to Syntenin 1 (Syndecan binding protein 1) (Melanoma), mRNA
 XM_376989 Homo sapiens similar to dJ54B20.4 (novel KRAB box containing C2H2 type 2), mRNA
 XM_377000 Homo sapiens hypothetical gene supported by AK096379 (LOC401589), mRNA
 XM_377002 Homo sapiens hypothetical gene supported by AK096379 (LOC401590), mRNA
 XM_377012 Homo sapiens similar to Spindlin-like protein 2 (SPIN-2) (LOC401591), mRNA
 XM_377014 Homo sapiens Cdc42 guanine nucleotide exchange factor (GEF) 9 (ARHGEF9), mRNA
 XM_377018 Homo sapiens LOC401594 (LOC401594), mRNA
 XM_377019 Homo sapiens LOC401595 (LOC401595), mRNA
 XM_377024 Homo sapiens similar to adaptor-related protein complex 2, beta 1 subunit; a (LOC401596), mRNA
 XM_377025 Homo sapiens similar to adaptor-related protein complex 2, beta 1 subunit; b (LOC401597), mRNA
 XM_377026 Homo sapiens similar to adaptor-related protein complex 2, beta 1 subunit; b (LOC401598), mRNA
 XM_377027 Homo sapiens similar to heat shock 70kD protein binding protein; progesterone (LOC401599), mRNA
 XM_377028 Homo sapiens similar to histone acetyltransferase (LOC401606), mRNA
 XM_377031 Homo sapiens similar to nuclear RNA export factor 2; TAP like protein 2 (LOC401607), mRNA
 XM_377032 Homo sapiens G protein-coupled receptor-associated sorting protein (GASP), mRNA
 XM_377033 Homo sapiens similar to STELLA (LOC401611), mRNA
 XM_377034 Homo sapiens similar to mitochondrial carrier triple repeat 1 (LOC401612), mRNA
 XM_377041 Homo sapiens similar to hypothetical protein MGC15416 (LOC401613), mRNA
 XM_377053 Homo sapiens hypothetical gene supported by BC040297 (LOC401619), mRNA
 XM_377060 Homo sapiens hypothetical protein LOC203547 (LOC203547), mRNA
 XM_377062 Homo sapiens similar to LINE-1 REVERSE TRANSCRIPTASE HOMOLOG (LOC401620), mRNA
 XM_377071 Homo sapiens similar to LINE-1 REVERSE TRANSCRIPTASE HOMOLOG (LOC401621), mRNA
 XM_377072 Homo sapiens similar to LINE-1 REVERSE TRANSCRIPTASE HOMOLOG (LOC401622), mRNA
 XM_377073 Homo sapiens similar to CXYorf1-related protein (LOC401624), mRNA
 XM_377076 Homo sapiens hypothetical gene supported by AK097803; BC017239 (LOC401625), mRNA
 XM_377087 Homo sapiens similar to hypothetical protein FLJ33610 (LOC401627), mRNA
 XM_377097 Homo sapiens similar to RNA binding motif protein, Y-linked, family 1 (LOC401628), mRNA
 XM_377098 Homo sapiens similar to Transcript Y 6 protein (LOC401633), mRNA
 XM_377102 Homo sapiens LOC401634 (LOC401634), mRNA
 XM_377104 Homo sapiens LOC401635 (LOC401635), mRNA
 XM_377109 Homo sapiens similar to 40S ribosomal protein SA (P40) (34/67 kDa laminin A), mRNA
 XM_377110 Homo sapiens similar to peptidylprolyl isomerase A (cyclophilin A) (LOC401640), mRNA
 XM_377115 Homo sapiens similar to PRED65 (LOC401641), mRNA
 XM_377117 Homo sapiens similar to ba182L21.1 (novel protein similar to hypothetical protein), mRNA
 XM_377122 Homo sapiens similar to Guanine nucleotide-binding protein G(i), alpha-2 subunit (LOC401642), mRNA
 XM_377129 Homo sapiens similar to golgi autoantigen, golgin subfamily a, 7 (LOC401643), mRNA
 XM_377133 Homo sapiens similar to deleted in malignant brain tumors 1; crp-ductin; vorticeptin (LOC401644), mRNA
 XM_377136 Homo sapiens similar to double homeobox protein (LOC401650), mRNA

XM_377137 Homo sapiens similar to double homeobox protein (LOC401851), mRNA
 XM_377140 Homo sapiens similar to double homeobox protein (LOC401852), mRNA
 XM_377142 Homo sapiens similar to double homeobox protein (LOC401853), mRNA
 XM_377143 Homo sapiens similar to double homeobox protein (LOC401854), mRNA
 XM_377147 Homo sapiens similar to RIKEN cDNA 1500011L16 (LOC401860), mRNA
 XM_377154 Homo sapiens similar to seven transmembrane helix receptor (LOC401861),
 XM_377155 Homo sapiens similar to Olfactory receptor 51H1 (LOC401863), mRNA
 XM_377156 Homo sapiens similar to Olfactory receptor 51T1 (LOC401865), mRNA
 XM_377158 Homo sapiens similar to Olfactory receptor 51A4 (LOC401866), mRNA
 XM_377159 Homo sapiens similar to Olfactory receptor 51A2 (LOC401867), mRNA
 XM_377179 Homo sapiens similar to seven transmembrane helix receptor (LOC401875),
 XM_377185 Homo sapiens similar to filamin-binding LIM protein-1; migfilin (LOC401879),
 XM_377189 Homo sapiens similar to Metabotropic glutamate receptor 5 precursor (mGluR)
 XM_377200 Homo sapiens similar to seven transmembrane helix receptor (LOC401887),
 XM_377218 Homo sapiens similar to MGC15937 protein (LOC401894), mRNA
 XM_377222 Homo sapiens similar to seven transmembrane helix receptor (LOC401896),
 XM_377230 Homo sapiens similar to U2af1 protein (LOC401703), mRNA
 XM_377231 Homo sapiens similar to heterogeneous nuclear ribonucleoprotein C isoform
 XM_377240 Homo sapiens similar to peptidylprolyl isomerase A (cyclophilin A) (LOC4017
 XM_377259 Homo sapiens similar to S-phase kinase-associated protein 1A isoform a; or
 XM_377262 Homo sapiens similar to Hypothetical protein CBG01854 (LOC401716), mR
 XM_377265 Homo sapiens similar to fidgetin (LOC401720), mRNA
 XM_377278 Homo sapiens similar to ribosomal protein L6 (LOC401725), mRNA
 XM_377283 Homo sapiens similar to Heat shock cognate 71 kDa protein (LOC401726), r
 XM_377285 Homo sapiens similar to 60S ribosomal protein L11 (LOC401727), mRNA
 XM_377287 Homo sapiens similar to MAL13P1.296 (LOC401728), mRNA
 XM_377296 Homo sapiens similar to Hypothetical protein CBG23155 (LOC401732), mR
 XM_377305 Homo sapiens similar to Hypothetical protein CBG01089 (LOC401740), mR
 XM_377306 Homo sapiens similar to vav-1 interacting Kruppel-like protein isoform b (LO
 XM_377337 Homo sapiens similar to cerebellin (LOC401766), mRNA
 XM_377338 Homo sapiens KIAA0323 (KIAA0323), mRNA
 XM_377343 Homo sapiens similar to hypothetical protein DJ667H12.2 (LOC401778), mR
 XM_377355 Homo sapiens KIAA0602 protein (KIAA0602), mRNA
 XM_377369 Homo sapiens similar to hypothetical protein FLJ36144 (LOC401806), mR/
 XM_377374 Homo sapiens similar to hypothetical protein FLJ36144 (LOC401808), mR/
 XM_377376 Homo sapiens similar to hypothetical protein FLJ36144 (LOC401811), mR/
 XM_377377 Homo sapiens similar to LRRGT00052 (LOC401812), mRNA
 XM_377383 Homo sapiens similar to stereoclin (LOC401815), mRNA
 XM_377388 Homo sapiens similar to antigen Cs44 (LOC401819), mRNA
 XM_377380 Homo sapiens similar to LOC375757 protein (LOC401820), mRNA
 XM_377384 Homo sapiens similar to melanoma chondroitin sulfate proteoglycan (LOC40
 XM_377385 Homo sapiens similar to MKI67 (FHA domain) interacting nuclear phospho
 XM_377407 Homo sapiens similar to hypothetical protein 4732467B22 (LOC401827), mR
 XM_377408 Homo sapiens similar to mast cell protease-11 (LOC401828), mRNA
 XM_377412 Homo sapiens similar to 3-phosphoinositide dependent protein kinase-1 (hPI
 XM_377414 Homo sapiens similar to Group X secretory phospholipase A2 precursor (Phc
 XM_377416 Homo sapiens similar to hypothetical protein MGC33867 (LOC401833), mR
 XM_377420 Homo sapiens similar to hypothetical protein BC011981 (LOC401837), mR
 XM_377423 Homo sapiens similar to Immunoglobulin heavy chain (LOC401841), mRNA
 XM_377424 Homo sapiens similar to Ig H-chain V-region (DP-40) (LOC401842), mRNA
 XM_377425 Homo sapiens similar to protein kinase related to Raf protein kinases; Metho
 XM_377426 Homo sapiens similar to IGHV gene product (LOC401845), mRNA
 XM_377429 Homo sapiens similar to IGHV gene product (LOC401846), mRNA
 XM_377439 Homo sapiens similar to hypothetical protein FLJ36144 (LOC401858), mR/
 XM_377444 Homo sapiens similar to peptidylprolyl isomerase A (cyclophilin A) (LOC401
 XM_377445 Homo sapiens similar to RIKEN cDNA C330003B14 (LOC401860), mRNA
 XM_377446 Homo sapiens similar to RIKEN cDNA C330003B14 (LOC401861), mRNA

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XM_377447 Homo sapiens similar to ribosomal protein S1 (LOC401862), mRNA
 XM_377451 Homo sapiens LOC401867 (LOC401867), mRNA
 XM_377455 Homo sapiens LOC401868 (LOC401868), mRNA
 XM_377457 Homo sapiens similar to 40S RIBOSOMAL PROTEIN S12 (LOC401870), mF
 XM_377464 Homo sapiens suppressor of Ty 6 homolog (S. cerevisiae) (SUPT6H), mRNA
 XM_377475 Homo sapiens similar to CDNA sequence BC004853 (LOC401883), mRNA
 XM_377476 Homo sapiens similar to Arf2-prov protein (LOC401884), mRNA
 XM_377477 Homo sapiens similar to 60S ribosomal protein L21 (LOC401885), mRNA
 XM_377480 Homo sapiens similar to 60S ribosomal protein L17 (L23) (Amino acid starva
 XM_377488 Homo sapiens similar to bA526D8.2 (novel protein similar to KIAA1074) (LO
 XM_377496 Homo sapiens similar to hypothetical protein (LOC401894), mRNA
 XM_377498 Homo sapiens KIAA0863 protein (KIAA0863), mRNA
 XM_377500 Homo sapiens similar to ribosomal protein S15; rat insulinoma gene (LOC40
 XM_377506 Homo sapiens egf-like module containing, mucin-like, hormone receptor-like
 XM_377511 Homo sapiens similar to 60S ribosomal protein L10 (QM protein homolog) (L
 XM_377512 Homo sapiens similar to glyceraldehyde 3-phosphate dehydrogenase (LOC4
 XM_377514 Homo sapiens similar to hypothetical protein FLJ38281 (LOC401898), mRN
 XM_377515 Homo sapiens similar to zinc finger protein 433 (LOC401899), mRNA
 XM_377516 Homo sapiens similar to ribosomal protein L28; 60S ribosomal protein L28 (L
 XM_377521 Homo sapiens similar to 60S ribosomal protein L23a (LOC401904), mRNA
 XM_377522 Homo sapiens similar to fibroblast growth factor receptor 3 (LOC401907), mF
 XM_377527 Homo sapiens similar to 60S ribosomal protein L29 (Cell surface heparin bin
 XM_377529 Homo sapiens similar to Zinc finger protein 345 (Zinc finger protein HZF10) (L
 XM_377533 Homo sapiens similar to Pregnancy-specific beta-1-glycoprotein 4 precursor
 XM_377537 Homo sapiens similar to OPA3 protein; Optic atrophy 3 (Iraqi-Jewish optic at
 XM_377538 Homo sapiens similar to Mucin 4 (Tracheobronchial mucin) (LOC401923), m
 XM_377553 Homo sapiens similar to zinc finger protein (LOC401932), mRNA
 XM_377554 Homo sapiens similar to hypothetical protein MGC4734 (LOC401933), mRN/
 XM_377555 Homo sapiens similar to RIKEN cDNA 5830442J12 (LOC401934), mRNA
 XM_377556 Homo sapiens similar to Hypothetical protein CBG06524 (LOC401936), mRN
 XM_377558 Homo sapiens similar to elongation factor 1 delta (LOC401937), mRNA
 XM_377563 Homo sapiens similar to Hypothetical protein DJ845O24.1 (LOC401938), mF
 XM_377565 Homo sapiens similar to Hypothetical protein DJ845O24.1 (LOC401939), mF
 XM_377566 Homo sapiens similar to dJ845O24.1 (Melanoma Preferentially Expressed A
 XM_377568 Homo sapiens similar to hypothetical protein (LOC401941), mRNA
 XM_377570 Homo sapiens similar to Hypothetical protein DJ845O24.1 (LOC401942), mF
 XM_377577 Homo sapiens hypothetical gene supported by AK127275 (LOC401944), mR
 XM_377579 Homo sapiens similar to cICK0721Q.2 (60S Ribosomal Protein L12 LIKE pro
 XM_377580 Homo sapiens similar to DC2 protein (LOC401946), mRNA
 XM_377583 Homo sapiens similar to ribosomal protein L27 (LOC401947), mRNA
 XM_377585 Homo sapiens similar to receptor tyrosine phosphatase (LOC401948), mRN/
 XM_377586 Homo sapiens similar to MGC52970 protein (LOC401949), mRNA
 XM_377593 Homo sapiens similar to dJ612B15.1 (novel protein similar to 60S ribosomal
 XM_377594 Homo sapiens similar to sporulation-induced transcript 4-associated protein :
 XM_377595 Homo sapiens similar to beta-actin (LOC401956), mRNA
 XM_377597 Homo sapiens similar to glyceraldehyde 3-phosphate dehydrogenase (LOC4
 XM_377599 Homo sapiens similar to hypothetical protein MGC8902 (LOC401962), mRN/
 XM_377600 Homo sapiens similar to hypothetical protein MGC8902 (LOC401963), mRN/
 XM_377601 Homo sapiens similar to AG3 (LOC401964), mRNA
 XM_377611 Homo sapiens similar to Olfactory receptor 10J6 (LOC401973), mRNA
 XM_377613 Homo sapiens similar to ribosomal protein S2; 40S ribosomal protein S2 (LO
 XM_377618 Homo sapiens LOC401976 (LOC401976), mRNA
 XM_377620 Homo sapiens similar to 14.5 kDa translational inhibitor protein (p14.5) (UK1
 XM_377630 Homo sapiens similar to RIKEN cDNA 2610020C11 (LOC401983), mRNA
 XM_377631 Homo sapiens similar to beta actin (LOC401987), mRNA
 XM_377633 Homo sapiens similar to Olfactory receptor 2G2 (LOC401990), mRNA
 XM_377635 Homo sapiens similar to CDC-like kinase 3; cdc2/CDC28-like protein kinase

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XM_377643 Homo sapiens similar to Olfactory receptor 2T2 (LOC401992), mRNA
 XM_377644 Homo sapiens similar to Olfactory receptor 2T5 (LOC401993), mRNA
 XM_377645 Homo sapiens similar to seven transmembrane helix receptor (LOC401994), mRNA
 XM_377649 Homo sapiens similar to 14.5 kDa translational inhibitor protein (p14.5) (UK1)
 XM_377655 Homo sapiens similar to seven transmembrane helix receptor (LOC401997), mRNA
 XM_377657 Homo sapiens similar to Olfactory receptor 2T5 (LOC401998), mRNA
 XM_377658 Homo sapiens similar to seven transmembrane helix receptor (LOC401999), mRNA
 XM_377659 Homo sapiens similar to Olfactory receptor 2T11 (LOC402000), mRNA
 XM_377660 Homo sapiens similar to seven transmembrane helix receptor (LOC402001), mRNA
 XM_377661 Homo sapiens similar to Olfactory receptor 5BF1 (LOC402002), mRNA
 XM_377662 Homo sapiens similar to Olfactory receptor 2T4 (LOC402003), mRNA
 XM_377663 Homo sapiens similar to Olfactory receptor 2T1 (Olfactory receptor 1-25) (OF)
 XM_377664 Homo sapiens similar to Olfactory receptor 2T2 (LOC402006), mRNA
 XM_377665 Homo sapiens similar to Olfactory receptor 2T3 (LOC402007), mRNA
 XM_377666 Homo sapiens similar to Olfactory receptor 2T5 (LOC402008), mRNA
 XM_377668 Homo sapiens similar to hypothetical protein F830045P16 (LOC402009), mRNA
 XM_377675 Homo sapiens similar to dJ1187J4.2 (novel protein similar to rat RYF3) (LOC)
 XM_377687 Homo sapiens similar to hypothetical protein FLJ21347 (LOC402027), mRNA
 XM_377690 Homo sapiens similar to extensin-like protein (LOC402030), mRNA
 XM_377691 Homo sapiens LOC402032 (LOC402032), mRNA
 XM_377694 Homo sapiens similar to LOC284861 protein (LOC402034), mRNA
 XM_377695 Homo sapiens similar to LOC284861 protein (LOC402035), mRNA
 XM_377696 Homo sapiens similar to carbonic anhydrase XV (LOC402036), mRNA
 XM_377700 Homo sapiens similar to LOC284861 protein (LOC402038), mRNA
 XM_377713 Homo sapiens similar to SRR1-like protein (LOC402055), mRNA
 XM_377715 Homo sapiens similar to Small nuclear ribonucleoprotein associated protein I
 XM_377716 Homo sapiens similar to 40S ribosomal protein S17 (LOC402057), mRNA
 XM_377717 Homo sapiens similar to dJ1119A7.3 (PUTATIVE novel protein similar to HP)
 XM_377720 Homo sapiens hypothetical protein BC012882 (LOC150356), mRNA
 XM_377721 Homo sapiens similar to Hypothetical protein CBG23588 (LOC402064), mRNA
 XM_377725 Homo sapiens similar to LWamide neuropeptide precursor protein (LOC40206)
 XM_377728 Homo sapiens similar to Hypothetical protein CBG08601 (LOC402067), mRNA
 XM_377732 Homo sapiens similar to egg envelope component ZPAX (LOC402068), mRNA
 XM_377741 Homo sapiens similar to SPCPB16A4.07c (LOC402072), mRNA
 XM_377742 Homo sapiens KIAA1940 protein (KIAA1940), mRNA
 XM_377751 Homo sapiens similar to Ig kappa chain V region (Z3) - human (LOC402089)
 XM_377754 Homo sapiens LOC402090 (LOC402090), mRNA
 XM_377755 Homo sapiens similar to hypothetical protein (LOC402094), mRNA
 XM_377756 Homo sapiens similar to dJ908M14.1.3 (ribosomal protein S21, isoform 3) (L)
 XM_377760 Homo sapiens similar to ribosomal protein L22 (LOC402098), mRNA
 XM_377761 Homo sapiens similar to ribosomal protein L22 (LOC402100), mRNA
 XM_377766 Homo sapiens similar to hypothetical protein DKFZp434P0315 (LOC402103)
 XM_377768 Homo sapiens similar to hypothetical protein FLJ10462 (LOC402104), mRNA
 XM_377771 Homo sapiens similar to 60S ribosomal protein L6 (TAX-responsive enhance
 XM_377774 Homo sapiens kinesin family member 5C (KIF5C), mRNA
 XM_377776 Homo sapiens similar to Acidic ribosomal phosphoprotein P0 (LOC402109), mRNA
 XM_377778 Homo sapiens LOC402110 (LOC402110), mRNA
 XM_377783 Homo sapiens similar to RIKEN cDNA A930041G11 gene (LOC402117), mRNA
 XM_377786 Homo sapiens similar to ribosomal protein L23 (LOC402120), mRNA
 XM_377797 Homo sapiens similar to laminin receptor-like protein LAMRL5 (LOC402123), mRNA
 XM_377803 Homo sapiens similar to RIKEN cDNA 1700112C13 (LOC402128), mRNA
 XM_377809 Homo sapiens similar to Olfactory receptor 5K2 (LOC402135), mRNA
 XM_377811 Homo sapiens similar to transcription factor INI (LOC402136), mRNA
 XM_377815 Homo sapiens similar to chromosome 11 open reading frame2; chromosome
 XM_377818 Homo sapiens similar to mesenchymal stem cell protein DSC92; neurite outg
 XM_377820 Homo sapiens similar to ribosomal protein L7-like 1 (LOC402152), mRNA
 XM_377823 Homo sapiens similar to p53 apoptosis effector related to Pmp22; p53 apopt

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XM_377824 Homo sapiens similar to Kinesin-like protein KIF3A (Microtubule plus end-din
 XM_377828 Homo sapiens similar to GLP_171_8870_6279 (LOC402160), mRNA
 XM_377829 Homo sapiens similar to hypothetical protein MGC45871 (LOC402161), mRN
 XM_377830 Homo sapiens similar to deubiquitinating enzyme 3 (LOC402164), mRNA
 XM_377831 Homo sapiens similar to deubiquitinating enzyme 3 (LOC402165), mRNA
 XM_377832 Homo sapiens similar to deubiquitinating enzyme 3 (LOC402166), mRNA
 XM_377834 Homo sapiens similar to deubiquitinating enzyme 3 (LOC402167), mRNA
 XM_377835 Homo sapiens similar to deubiquitinating enzyme 3 (LOC402168), mRNA
 XM_377836 Homo sapiens similar to deubiquitinating enzyme 3 (LOC402169), mRNA
 XM_377837 Homo sapiens similar to deubiquitinating enzyme 3 (LOC402170), mRNA
 XM_377841 Homo sapiens similar to 60S ribosomal protein L21 (LOC402176), mRNA
 XM_377845 Homo sapiens similar to ribosomal protein S21; 40S ribosomal protein S21 (I
 XM_377847 Homo sapiens similar to ras-related C3 botulinum toxin substrate 1 isoform F
 XM_377849 Homo sapiens similar to alanyl tna synthetase (LOC402188), mRNA
 XM_377861 Homo sapiens similar to glycine-rich protein (LOC402194), mRNA
 XM_377875 Homo sapiens similar to Transcription initiation factor TFIID 28 kDa subunit (I
 XM_377877 Homo sapiens similar to Transcription initiation factor TFIID 28 kDa subunit (I
 XM_377878 Homo sapiens similar to Transcription initiation factor TFIID 28 kDa subunit (I
 XM_377879 Homo sapiens similar to Transcription initiation factor TFIID 28 kDa subunit (I
 XM_377880 Homo sapiens similar to Transcription initiation factor TFIID 28 kDa subunit (I
 XM_377881 Homo sapiens similar to Transcription initiation factor TFIID 28 kDa subunit (I
 XM_377882 Homo sapiens similar to Transcription initiation factor TFIID 28 kDa subunit (I
 XM_377883 Homo sapiens similar to Transcription initiation factor TFIID 28 kDa subunit (I
 XM_377884 Homo sapiens similar to Transcription initiation factor TFIID 28 kDa subunit (I
 XM_377885 Homo sapiens similar to Transcription initiation factor TFIID 28 kDa subunit (I
 XM_377886 Homo sapiens similar to Transcription initiation factor TFIID 28 kDa subunit (I
 XM_377887 Homo sapiens similar to Transcription initiation factor TFIID 28 kDa subunit (I
 XM_377889 Homo sapiens similar to Transcription initiation factor TFIID 28 kDa subunit (I
 XM_377896 Homo sapiens similar to ribosomal protein L13a; 60S ribosomal protein L13a
 XM_377904 Homo sapiens similar to cytoplasmic beta-actin (LOC402218), mRNA
 XM_377907 Homo sapiens similar to beta-glucuronidase (LOC402223), mRNA
 XM_377910 Homo sapiens similar to RIKEN cDNA A730017C20 (LOC402228), mRNA
 XM_377911 Homo sapiens similar to hypothetical protein E230025N22 (LOC402231), mF
 XM_377912 Homo sapiens KIAA0194 protein (KIAA0194), mRNA
 XM_377918 Homo sapiens similar to Ac1147 (LOC402235), mRNA
 XM_377919 Homo sapiens similar to RIKEN cDNA 2310040C09 (LOC402237), mRNA
 XM_377920 Homo sapiens similar to Selenophosphate synthetase 1 (LOC402238), mRN
 XM_377924 Homo sapiens similar to Olfactory receptor 4F3 (LOC402242), mRNA
 XM_377925 Homo sapiens similar to hypothetical protein FLJ37300 (LOC402244), mRN/
 XM_377926 Homo sapiens similar to mesenchymal stem cell protein DSC92; neurite outg
 XM_377927 Homo sapiens similar to olfactory receptor MOR267-3 (LOC402246), mRNA
 XM_377928 Homo sapiens similar to Calgizzarin (S100C protein) (MLN 70) (LOC402247)
 XM_377929 Homo sapiens similar to ribosomal protein L31 (LOC402248), mRNA
 XM_377931 Homo sapiens similar to olfactory receptor MOR145-2 (LOC402249), mRNA
 XM_377932 Homo sapiens similar to 4930579E17Rik protein (LOC402250), mRNA
 XM_377933 Homo sapiens similar to MGC76216 protein (LOC402251), mRNA
 XM_377934 Homo sapiens similar to peptidylprolyl isomerase A (LOC402252), mRNA
 XM_377935 Homo sapiens similar to FLJ40113 protein (LOC402253), mRNA
 XM_377938 Homo sapiens similar to ELK1 (LOC402257), mRNA
 XM_377941 Homo sapiens similar to equilibrative nucleoside transporter 4; hENT4 (LOC/
 XM_377942 Homo sapiens similar to vomeronasal receptor V1RC3 (LOC402273), mRNA
 XM_377943 Homo sapiens similar to Opioid binding protein/cell adhesion molecule precu
 XM_377944 Homo sapiens similar to protein kinase related to Raf protein kinases; Melho
 XM_377945 Homo sapiens similar to metabotropic glutamate receptor 8; G protein-coupl
 XM_377946 Homo sapiens similar to GTF2I repeat domain containing 1 isoform 2; Williar
 XM_377947 Homo sapiens similar to opposite strand transcription unit to Stag3; Gats pro
 XM_377949 Homo sapiens similar to GrpE protein homolog 1, mitochondrial precursor (V

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XM_377950 Homo sapiens similar to GTF2I repeat domain containing 1 isoform 2; Williar
 XM_377951 Homo sapiens similar to peptidylprolyl isomerase A (LOC402284), mRNA
 XM_377955 Homo sapiens hypothetical protein DKFZP434A0225 (DKFZP434A0225), mf
 XM_377956 Homo sapiens similar to RIKEN cDNA 4930511M11 (LOC402286), mRNA
 XM_377957 Homo sapiens similar to ribosomal protein S3a; 40S ribosomal protein S3a; \
 XM_377958 Homo sapiens similar to Ser/Thr protein kinase PAR-1Balpha (LOC402289),
 XM_377959 Homo sapiens similar to Nuclear protein Hcc-1 (HSPC316) (Proliferation ass
 XM_377961 Homo sapiens similar to opposite strand transcription unit to Stag3; Gats pro
 XM_377962 Homo sapiens postmeiotic segregation increased 2-like 1 (PMS2L1), mRNA
 XM_377964 Homo sapiens similar to 40S RIBOSOMAL PROTEIN SA (P40) (3467 KD L
 XM_377969 Homo sapiens similar to 60S ribosomal protein L23a (LOC402294), mRNA
 XM_377970 Homo sapiens similar to Argininosuccinate synthase (Citrulline--aspartate lig.
 XM_377972 Homo sapiens similar to Triosephosphate isomerase (TIM) (LOC402298), ml
 XM_377973 Homo sapiens similar to aldo-keto reductase family 1, member B10; aldose r
 XM_377974 Homo sapiens similar to beta-tubulin (LOC402300), mRNA
 XM_377976 Homo sapiens similar to Nucleoside diphosphate kinase, mitochondrial prec
 XM_377995 Homo sapiens similar to Olfactory receptor 2A1 (LOC402317), mRNA
 XM_377997 Homo sapiens similar to 60S ribosomal protein L32 (LOC402318), mRNA
 XM_377998 Homo sapiens similar to Huntingtin interacting protein K (LOC402319), mRN
 XM_377999 Homo sapiens similar to BET1 homolog (Golgi vesicular membrane traffickin
 XM_378001 Homo sapiens similar to TSH receptor suppressor element-binding protein-1
 XM_378002 Homo sapiens similar to ppg3 (LOC402322), mRNA
 XM_378003 Homo sapiens similar to S-adenosylmethionine decarboxylase 1; S-adenosyl
 XM_378007 Homo sapiens similar to hypothetical protein PFL1865w (LOC402324), mRN
 XM_378008 Homo sapiens similar to ENSANGP00000017949 (LOC402325), mRNA
 XM_378009 Homo sapiens similar to hypothetical protein (LOC402326), mRNA
 XM_378010 Homo sapiens similar to stage-specific S antigen homolog (LOC402327), mF
 XM_378014 Homo sapiens similar to ubiquitin-specific protease 17-like protein (LOC4022
 XM_378015 Homo sapiens similar to chromosome 11 open reading frame2; chromosome
 XM_378018 Homo sapiens similar to chromosome 11 open reading frame2; chromosome
 XM_378028 Homo sapiens similar to L21 ribosomal protein (LOC402336), mRNA
 XM_378031 Homo sapiens similar to short chain dehydrogenase reductase 9 (LOC40233
 XM_378033 Homo sapiens similar to chromosome 20 open reading frame 6 (LOC402340
 XM_378035 Homo sapiens similar to fatty acid binding protein 9, testis; testis lipid bindin
 XM_378036 Homo sapiens similar to tropomyosin 3, gamma (LOC402344), mRNA
 XM_378043 Homo sapiens similar to RIKEN cDNA 1700091F14 (LOC402353), mRNA
 XM_378044 Homo sapiens similar to hypothetical protein (LOC402354), mRNA
 XM_378046 Homo sapiens similar to RIKEN cDNA 1700091F14 (LOC402355), mRNA
 XM_378052 Homo sapiens similar to Interferon omega-1 precursor (Interferon alpha-II-1)
 XM_378054 Homo sapiens similar to hypothetical protein (LOC402360), mRNA
 XM_378062 Homo sapiens similar to 6-pyruvoyl-tetrahydropterin synthase (LOC402365),
 XM_378063 Homo sapiens similar to 6-pyruvoyl-tetrahydropterin synthase (LOC402366),
 XM_378064 Homo sapiens LOC402367 (LOC402367), mRNA
 XM_378067 Homo sapiens similar to phosphoglucosyltransferase 5 (LOC402368), mRNA
 XM_378078 Homo sapiens chromosome 9 open reading frame 4 (C9orf4), mRNA
 XM_378080 Homo sapiens similar to beta-1,3-N-acetylglucosaminyltransferase 5 (LOC4C
 XM_378087 Homo sapiens similar to ligand-independent activating molecule for estrogen
 XM_378089 Homo sapiens similar to LOC286220 protein (LOC402381), mRNA
 XM_378090 Homo sapiens similar to F4N2.10 (LOC402382), mRNA
 XM_378102 Homo sapiens similar to ENSANGP00000002367 (LOC402386), mRNA
 XM_378103 Homo sapiens similar to hydroxymethylpterin pyrophosphokinase-dihydropte
 XM_378113 Homo sapiens similar to LOC142827 protein (LOC402402), mRNA
 XM_378116 Homo sapiens similar to DKFZP434C0047 protein (LOC402403), mRNA
 XM_378118 Homo sapiens similar to PAGE-5 protein (LOC402404), mRNA
 XM_378123 Homo sapiens similar to Mitochondrial import receptor subunit TOM20 homol
 XM_378124 Homo sapiens similar to bA203116.1 (KIAA0970 protein) (LOC402414), mRN
 XM_378125 Homo sapiens similar to hypothetical protein MGC57211 (LOC402415), mRN

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XM_378128 Homo sapiens similar to dJ19N1.1 (novel protein) (LOC402416), mRNA
 XM_378137 Homo sapiens similar to olfactory receptor MOR262-9 (LOC402424), mRNA
 XM_378143 Homo sapiens similar to tau tubulin Kinase 2; tau-tubulin kinase (LOC402425)
 XM_378152 Homo sapiens similar to BA203116.1 (KIAA0970 protein) (LOC402432), mRNA
 XM_378155 Homo sapiens similar to hydroxymethylpterin pyrophosphokinase-dihydropterin
 XM_378156 Homo sapiens similar to trophinin; melanoma antigen, family D, 3; trophinin-2
 XM_378158 Homo sapiens similar to hypothetical protein FLJ90430 (LOC402437), mRNA
 XM_378172 Homo sapiens spermatogenesis-related protein 8 (MGC44294), mRNA
 XM_378173 Homo sapiens hypothetical protein MGC18216 (MGC18216), mRNA
 XM_378175 Homo sapiens hypothetical protein BC017488 (LOC124446), mRNA
 XM_378177 Homo sapiens hypothetical gene supported by NM_078471 (LOC399700), m
 XM_378178 Homo sapiens hypothetical protein MGC9913 (MGC9913), mRNA
 XM_378180 Homo sapiens hypothetical protein MGC10812 (MGC10812), mRNA
 XM_378181 Homo sapiens KIAA1041 protein (KIAA1041), mRNA
 XM_378183 Homo sapiens hypothetical protein MGC5457 (MGC5457), mRNA
 XM_378184 Homo sapiens KIAA1383 protein (KIAA1383), mRNA
 XM_378185 Homo sapiens hypothetical gene supported by AL833273; NM_014644 (LOC
 XM_378186 Homo sapiens hypothetical protein MGC15634 (MGC15634), mRNA
 XM_378187 Homo sapiens hypothetical protein MGC4473 (MGC4473), mRNA
 XM_378189 Homo sapiens hypothetical protein MGC15705 (MGC15705), mRNA
 XM_378190 Homo sapiens hypothetical protein MGC10955 (MGC10955), mRNA
 XM_378191 Homo sapiens hypothetical protein PRO2964 (PRO2964), mRNA
 XM_378192 Homo sapiens hypothetical gene supported by NM_015583 (LOC399702), m
 XM_378193 Homo sapiens hypothetical protein MGC10981 (MGC10981), mRNA
 XM_378194 Homo sapiens hypothetical gene supported by NM_020669 (LOC399703), m
 XM_378195 Homo sapiens hypothetical gene supported by NM_001517 (LOC399704), m
 XM_378196 Homo sapiens hypothetical protein FLJ36112 (FLJ36112), mRNA
 XM_378197 Homo sapiens hypothetical protein FLJ14464 (FLJ14464), mRNA
 XM_378199 Homo sapiens hypothetical protein FLJ10232 (FLJ10232), mRNA
 XM_378200 Homo sapiens hypothetical gene supported by AK097673 (LOC399706), mR
 XM_378201 Homo sapiens hypothetical protein LOC282980 (LOC282980), mRNA
 XM_378202 Homo sapiens hypothetical gene supported by AK056101 (LOC399707), mR
 XM_378203 Homo sapiens hypothetical gene supported by BC055423 (LOC399708), mR
 XM_378207 Homo sapiens hypothetical protein LOC338588 (LOC338588), mRNA
 XM_378208 Homo sapiens hypothetical gene supported by AK125014 (LOC399713), mR
 XM_378210 Homo sapiens hypothetical gene supported by AK128810 (LOC399717), mR
 XM_378211 Homo sapiens hypothetical protein LOC254312 (LOC254312), mRNA
 XM_378215 Homo sapiens hypothetical gene supported by BC040880 (LOC399726), mR
 XM_378218 Homo sapiens hypothetical gene supported by BX537934 (LOC399736), mR
 XM_378219 Homo sapiens LOC399737 (LOC399737), mRNA
 XM_378223 Homo sapiens hypothetical gene supported by AK083334; AL833330; BC021
 XM_378224 Homo sapiens hypothetical gene supported by X06747; BC012158; NM_002
 XM_378226 Homo sapiens hypothetical protein LOC170371 (LOC170371), mRNA
 XM_378227 Homo sapiens hypothetical gene supported by AK083334; AL833330; BC021
 XM_378228 Homo sapiens hypothetical gene supported by AK083334; AL833330; BC021
 XM_378230 Homo sapiens hypothetical protein LOC283025 (LOC283025), mRNA
 XM_378232 Homo sapiens hypothetical protein LOC219690 (LOC219690), mRNA
 XM_378235 Homo sapiens LOC399785 (LOC399785), mRNA
 XM_378236 Homo sapiens hypothetical gene supported by AK091031 (LOC399786), mR
 XM_378238 Homo sapiens hypothetical protein LOC283050 (LOC283050), mRNA
 XM_378239 Homo sapiens hypothetical gene supported by AK056520; NM_153030 (LOC
 XM_378240 Homo sapiens hypothetical protein LOC170425 (LOC170425), mRNA
 XM_378247 Homo sapiens hypothetical gene supported by AK123344 (LOC399806), mR
 XM_378250 Homo sapiens hypothetical protein LOC92482 (LOC92482), mRNA
 XM_378251 Homo sapiens hypothetical protein LOC143188 (LOC143188), mRNA
 XM_378255 Homo sapiens hypothetical gene supported by AK128177 (LOC399827), mR
 XM_378257 Homo sapiens hypothetical gene supported by AK125849 (LOC399829), mR

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XM_378259 Homo sapiens hypothetical gene supported by BX647230 (LOC399832), mRNA
 XM_378260 Homo sapiens hypothetical gene supported by AK126615 (LOC399833), mRNA
 XM_378261 Homo sapiens hypothetical gene supported by BC004945; BC020442; BC06
 XM_378262 Homo sapiens hypothetical protein LOC284701 (LOC284701), mRNA
 XM_378266 Homo sapiens hypothetical gene supported by AY129010 (LOC399851), mRNA
 XM_378271 Homo sapiens hypothetical gene supported by AK002039 (LOC399865), mRNA
 XM_378272 Homo sapiens hypothetical gene supported by BX647519 (LOC399866), mRNA
 XM_378273 Homo sapiens hypothetical gene supported by AK127718 (LOC399867), mRNA
 XM_378276 Homo sapiens LOC399872 (LOC399872), mRNA
 XM_378277 Homo sapiens hypothetical gene supported by AK096475 (LOC399873), mRNA
 XM_378279 Homo sapiens hypothetical gene supported by BC040220 (LOC399875), mRNA
 XM_378280 Homo sapiens hypothetical gene supported by AK023501 (LOC399876), mRNA
 XM_378283 Homo sapiens hypothetical gene supported by AK127155 (LOC399879), mRNA
 XM_378286 Homo sapiens hypothetical gene supported by AK093366 (LOC399884), mRNA
 XM_378288 Homo sapiens hypothetical gene supported by AK123417 (LOC399886), mRNA
 XM_378297 Homo sapiens hypothetical gene supported by X15675 (LOC399912), mRNA
 XM_378299 Homo sapiens hypothetical gene supported by AK094674 (LOC399919), mRNA
 XM_378300 Homo sapiens hypothetical gene supported by BC039105 (LOC399920), mRNA
 XM_378301 Homo sapiens hypothetical gene supported by AL832797 (LOC399924), mRNA
 XM_378303 Homo sapiens hypothetical protein LOC283214 (LOC283214), mRNA
 XM_378304 Homo sapiens hypothetical gene supported by BC026292 (LOC399930), mRNA
 XM_378305 Homo sapiens LOC399933 (LOC399933), mRNA
 XM_378308 Homo sapiens LOC399945 (LOC399945), mRNA
 XM_378309 Homo sapiens LOC399951 (LOC399951), mRNA
 XM_378311 Homo sapiens hypothetical gene supported by AK124988 (LOC399954), mRNA
 XM_378312 Homo sapiens hypothetical protein LOC283143 (LOC283143), mRNA
 XM_378313 Homo sapiens LOC399955 (LOC399955), mRNA
 XM_378314 Homo sapiens hypothetical protein LOC283152 (LOC283152), mRNA
 XM_378316 Homo sapiens hypothetical gene supported by BX647608 (LOC399959), mRNA
 XM_378317 Homo sapiens LOC399961 (LOC399961), mRNA
 XM_378320 Homo sapiens hypothetical gene supported by AK125355 (LOC399971), mRNA
 XM_378321 Homo sapiens hypothetical gene supported by AK096370 (LOC399972), mRNA
 XM_378325 Homo sapiens hypothetical gene supported by BC031979 (LOC399978), mRNA
 XM_378326 Homo sapiens hypothetical gene supported by AK127362 (LOC399980), mRNA
 XM_378327 Homo sapiens hypothetical protein LOC283177 (LOC283177), mRNA
 XM_378328 Homo sapiens hypothetical gene supported by BC039168 (LOC399982), mRNA
 XM_378329 Homo sapiens hypothetical gene supported by AK090616 (LOC399983), mRNA
 XM_378330 Homo sapiens hypothetical gene supported by AK057909 (LOC399984), mRNA
 XM_378331 Homo sapiens hypothetical gene supported by AK056228 (LOC399986), mRNA
 XM_378332 Homo sapiens LOC399987 (LOC399987), mRNA
 XM_378336 Homo sapiens LOC399993 (LOC399993), mRNA
 XM_378339 Homo sapiens hypothetical gene supported by AK128230 (LOC400002), mRNA
 XM_378340 Homo sapiens LOC400004 (LOC400004), mRNA
 XM_378342 Homo sapiens hypothetical gene supported by BC003510; NM_002823 (LOC
 XM_378343 Homo sapiens hypothetical gene supported by AK097936 (LOC400007), mRNA
 XM_378344 Homo sapiens LOC400015 (LOC400015), mRNA
 XM_378346 Homo sapiens hypothetical protein LOC196415 (LOC196415), mRNA
 XM_378349 Homo sapiens LOC400019 (LOC400019), mRNA
 XM_378350 Homo sapiens hypothetical gene supported by BC047417 (LOC400027), mRNA
 XM_378353 Homo sapiens LOC400030 (LOC400030), mRNA
 XM_378354 Homo sapiens LOC400031 (LOC400031), mRNA
 XM_378355 Homo sapiens hypothetical protein LOC283332 (LOC283332), mRNA
 XM_378356 Homo sapiens hypothetical protein LOC283400 (LOC283400), mRNA
 XM_378358 Homo sapiens hypothetical gene supported by AK123741 (LOC400041), mRNA
 XM_378360 Homo sapiens hypothetical gene supported by BC009385 (LOC400043), mRNA
 XM_378362 Homo sapiens hypothetical gene supported by AK123272 (LOC400046), mRNA
 XM_378363 Homo sapiens LOC400047 (LOC400047), mRNA

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XM_378365 Homo sapiens LOC400050 (LOC400050), mRNA
 XM_378366 Homo sapiens hypothetical gene supported by AK124066 (LOC400051), mR
 XM_378367 Homo sapiens LOC400053 (LOC400053), mRNA
 XM_378368 Homo sapiens hypothetical protein LOC283392 (LOC283392), mRNA
 XM_378371 Homo sapiens hypothetical protein LOC338758 (LOC338758), mRNA
 XM_378372 Homo sapiens hypothetical protein LOC256021 (LOC256021), mRNA
 XM_378374 Homo sapiens hypothetical protein LOC338809 (LOC338809), mRNA
 XM_378379 Homo sapiens hypothetical protein LOC283432 (LOC283432), mRNA
 XM_378381 Homo sapiens hypothetical gene supported by BX648662 (LOC400070), mR
 XM_378388 Homo sapiens hypothetical protein LOC144742 (LOC144742), mRNA
 XM_378389 Homo sapiens hypothetical gene supported by AK057632; AL137270; BC05
 XM_378390 Homo sapiens hypothetical protein LOC144678 (LOC144678), mRNA
 XM_378392 Homo sapiens hypothetical gene supported by AK094824 (LOC400087), mR
 XM_378393 Homo sapiens hypothetical gene supported by AK126855 (LOC400088), mR
 XM_378394 Homo sapiens hypothetical protein LOC116437 (LOC116437), mRNA
 XM_378398 Homo sapiens LOC400092 (LOC400092), mRNA
 XM_378399 Homo sapiens hypothetical gene supported by BC024195 (LOC400099), mR
 XM_378404 Homo sapiens hypothetical gene supported by AK098387 (LOC400108), mR
 XM_378405 Homo sapiens LOC400111 (LOC400111), mRNA
 XM_378407 Homo sapiens hypothetical gene supported by BX648491 (LOC400115), mR
 XM_378411 Homo sapiens hypothetical gene supported by AK124383 (LOC400123), mR
 XM_378412 Homo sapiens LOC400125 (LOC400125), mRNA
 XM_378413 Homo sapiens hypothetical gene supported by BC025370 (LOC400128), mR
 XM_378414 Homo sapiens hypothetical gene supported by AF529010 (LOC400131), mR
 XM_378416 Homo sapiens LOC400134 (LOC400134), mRNA
 XM_378419 Homo sapiens hypothetical protein LOC144766 (LOC144766), mRNA
 XM_378421 Homo sapiens hypothetical gene supported by AK056689 (LOC400144), mR
 XM_378425 Homo sapiens LOC400151 (LOC400151), mRNA
 XM_378428 Homo sapiens hypothetical gene supported by BC035106 (LOC400154), mR
 XM_378430 Homo sapiens hypothetical protein LOC283480 (LOC283480), mRNA
 XM_378431 Homo sapiens hypothetical protein LOC283483 (LOC283483), mRNA
 XM_378434 Homo sapiens hypothetical gene supported by BC038751 (LOC400161), mR
 XM_378436 Homo sapiens hypothetical gene supported by BC034786 (LOC400163), mR
 XM_378437 Homo sapiens hypothetical gene supported by BX649107 (LOC400164), mR
 XM_378439 Homo sapiens hypothetical gene supported by BC041346 (LOC400167), mR
 XM_378441 Homo sapiens LOC400171 (LOC400171), mRNA
 XM_378449 Homo sapiens LOC400201 (LOC400201), mRNA
 XM_378452 Homo sapiens hypothetical protein LOC253970 (LOC253970), mRNA
 XM_378453 Homo sapiens hypothetical gene supported by AK125955 (LOC400208), mR
 XM_378454 Homo sapiens hypothetical protein LOC283547 (LOC283547), mRNA
 XM_378455 Homo sapiens hypothetical protein LOC283551 (LOC283551), mRNA
 XM_378456 Homo sapiens hypothetical gene supported by AK127576 (LOC400212), mR
 XM_378457 Homo sapiens hypothetical gene supported by BC055421 (LOC400213), mR
 XM_378460 Homo sapiens hypothetical gene supported by BC037850 (LOC400216), mR
 XM_378462 Homo sapiens hypothetical gene supported by AK026100 (LOC400221), mR
 XM_378465 Homo sapiens LOC400228 (LOC400228), mRNA
 XM_378467 Homo sapiens LOC400231 (LOC400231), mRNA
 XM_378470 Homo sapiens hypothetical gene supported by BC029835 (LOC400234), mR
 XM_378472 Homo sapiens LOC400236 (LOC400236), mRNA
 XM_378473 Homo sapiens hypothetical gene supported by AK093266 (LOC400238), mR
 XM_378476 Homo sapiens hypothetical gene supported by AK127179 (LOC400242), mR
 XM_378477 Homo sapiens LOC400243 (LOC400243), mRNA
 XM_378482 Homo sapiens LOC400249 (LOC400249), mRNA
 XM_378487 Homo sapiens hypothetical protein LOC145216 (LOC145216), mRNA
 XM_378488 Homo sapiens LOC400256 (LOC400256), mRNA
 XM_378489 Homo sapiens hypothetical gene supported by AK091459 (LOC400257), mR
 XM_378490 Homo sapiens hypothetical gene supported by AK127521 (LOC400262), mR

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XM_378491 Homo sapiens hypothetical gene supported by BC033241 (LOC400263), mRNA
 XM_378493 Homo sapiens hypothetical gene supported by AK127783 (LOC400302), mRNA
 XM_378494 Homo sapiens LOC400307 (LOC400307), mRNA
 XM_378496 Homo sapiens hypothetical gene supported by BC047459 (LOC400314), mRNA
 XM_378506 Homo sapiens hypothetical gene supported by AK125576; AL117445; BC03:
 XM_378507 Homo sapiens hypothetical protein LOC145845 (LOC145845), mRNA
 XM_378511 Homo sapiens hypothetical gene supported by BC031266 (LOC400368), mRNA
 XM_378512 Homo sapiens hypothetical gene supported by BX537772 (LOC400369), mRNA
 XM_378514 Homo sapiens hypothetical protein LOC283663 (LOC283663), mRNA
 XM_378515 Homo sapiens hypothetical gene supported by AK091917 (LOC400377), mRNA
 XM_378516 Homo sapiens hypothetical protein LOC255177 (LOC255177), mRNA
 XM_378517 Homo sapiens hypothetical protein MGC15885 (MGC15885), mRNA
 XM_378522 Homo sapiens hypothetical gene supported by BC043587 (LOC400386), mRNA
 XM_378523 Homo sapiens LOC400388 (LOC400388), mRNA
 XM_378525 Homo sapiens hypothetical protein LOC283731 (LOC283731), mRNA
 XM_378526 Homo sapiens LOC400393 (LOC400393), mRNA
 XM_378528 Homo sapiens LOC400398 (LOC400398), mRNA
 XM_378529 Homo sapiens hypothetical gene supported by AK022116 (LOC400400), mRNA
 XM_378532 Homo sapiens hypothetical protein LOC253044 (LOC253044), mRNA
 XM_378535 Homo sapiens LOC400411 (LOC400411), mRNA
 XM_378538 Homo sapiens hypothetical gene supported by AL137524 (LOC400433), mRNA
 XM_378542 Homo sapiens hypothetical protein LOC283761 (LOC283761), mRNA
 XM_378544 Homo sapiens hypothetical protein LOC283682 (LOC283682), mRNA
 XM_378545 Homo sapiens hypothetical gene supported by BC040875 (LOC400456), mRNA
 XM_378546 Homo sapiens hypothetical protein LOC145820 (LOC145820), mRNA
 XM_378549 Homo sapiens hypothetical protein LOC91948 (LOC91948), mRNA
 XM_378550 Homo sapiens hypothetical protein LOC145757 (LOC145757), mRNA
 XM_378551 Homo sapiens LOC400463 (LOC400463), mRNA
 XM_378553 Homo sapiens hypothetical gene supported by BC041891 (LOC400475), mRNA
 XM_378558 Homo sapiens hypothetical protein LOC146443 (LOC146443), mRNA
 XM_378562 Homo sapiens hypothetical gene supported by AL162011 (LOC400496), mRNA
 XM_378564 Homo sapiens LOC400500 (LOC400500), mRNA
 XM_378567 Homo sapiens hypothetical gene supported by BX640722 (LOC400505), mRNA
 XM_378571 Homo sapiens hypothetical gene supported by AK127191 (LOC400511), mRNA
 XM_378573 Homo sapiens hypothetical gene supported by AK025061 (LOC400512), mRNA
 XM_378576 Homo sapiens hypothetical gene supported by AK126852 (LOC400516), mRNA
 XM_378577 Homo sapiens hypothetical gene supported by AK123554 (LOC400517), mRNA
 XM_378578 Homo sapiens hypothetical gene supported by BC023258 (LOC400518), mRNA
 XM_378579 Homo sapiens hypothetical gene supported by AK097527 (LOC400522), mRNA
 XM_378582 Homo sapiens LOC400523 (LOC400523), mRNA
 XM_378586 Homo sapiens LOC400531 (LOC400531), mRNA
 XM_378588 Homo sapiens hypothetical gene supported by BC047414 (LOC400532), mRNA
 XM_378589 Homo sapiens hypothetical protein LOC283914 (LOC283914), mRNA
 XM_378590 Homo sapiens hypothetical gene supported by AK129756 (LOC400533), mRNA
 XM_378592 Homo sapiens LOC400534 (LOC400534), mRNA
 XM_378594 Homo sapiens hypothetical gene supported by AK128747 (LOC400535), mRNA
 XM_378595 Homo sapiens hypothetical gene supported by AK057373 (LOC400536), mRNA
 XM_378599 Homo sapiens hypothetical protein LOC283854 (LOC283854), mRNA
 XM_378601 Homo sapiens hypothetical gene supported by AK057319 (LOC400538), mRNA
 XM_378606 Homo sapiens hypothetical protein LOC283867 (LOC283867), mRNA
 XM_378607 Homo sapiens hypothetical gene supported by AL080152 (LOC400540), mRNA
 XM_378608 Homo sapiens hypothetical gene supported by AK096068 (LOC400541), mRNA
 XM_378609 Homo sapiens hypothetical gene supported by BC064480 (LOC400544), mRNA
 XM_378610 Homo sapiens LOC400545 (LOC400545), mRNA
 XM_378617 Homo sapiens hypothetical gene supported by BC040618 (LOC400548), mRNA
 XM_378620 Homo sapiens hypothetical gene supported by AK091834 (LOC400550), mRNA
 XM_378621 Homo sapiens hypothetical gene supported by AK125749 (LOC400551), mRNA

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XM_378623 Homo sapiens hypothetical gene supported by AK055320 (LOC400552), mRNA
 XM_378625 Homo sapiens hypothetical gene supported by AK126852 (LOC400553), mRNA
 XM_378626 Homo sapiens hypothetical gene supported by AK123554 (LOC400554), mRNA
 XM_378628 Homo sapiens hypothetical protein MGC23284 (MGC23284), mRNA
 XM_378629 Homo sapiens hypothetical gene supported by AK127064 (LOC400556), mRNA
 XM_378630 Homo sapiens hypothetical gene supported by AK055272 (LOC400557), mRNA
 XM_378631 Homo sapiens hypothetical gene supported by AK130578 (LOC400558), mRNA
 XM_378632 Homo sapiens LOC400560 (LOC400560), mRNA
 XM_378633 Homo sapiens LOC400561 (LOC400561), mRNA
 XM_378634 Homo sapiens LOC400562 (LOC400562), mRNA
 XM_378637 Homo sapiens LOC400565 (LOC400565), mRNA
 XM_378639 Homo sapiens hypothetical gene supported by AK093301 (LOC400567), mRNA
 XM_378642 Homo sapiens hypothetical protein LOC284009 (LOC284009), mRNA
 XM_378643 Homo sapiens hypothetical gene supported by BC043554 (LOC400568), mRNA
 XM_378646 Homo sapiens LOC400571 (LOC400571), mRNA
 XM_378648 Homo sapiens LOC400572 (LOC400572), mRNA
 XM_378649 Homo sapiens hypothetical gene supported by BC015790; BC041634 (LOC400573), mRNA
 XM_378650 Homo sapiens hypothetical gene supported by BC017752 (LOC400575), mRNA
 XM_378653 Homo sapiens hypothetical gene supported by AK098898 (LOC400577), mRNA
 XM_378655 Homo sapiens hypothetical protein LOC98597 (LOC98597), mRNA
 XM_378656 Homo sapiens hypothetical gene supported by AK093253 (LOC400579), mRNA
 XM_378660 Homo sapiens hypothetical gene supported by AK093253 (LOC400584), mRNA
 XM_378661 Homo sapiens hypothetical protein LOC339263 (LOC339263), mRNA
 XM_378664 Homo sapiens hypothetical gene supported by AK124344 (LOC400588), mRNA
 XM_378667 Homo sapiens hypothetical protein LOC147004 (LOC147004), mRNA
 XM_378668 Homo sapiens hypothetical gene supported by AK125932 (LOC400593), mRNA
 XM_378675 Homo sapiens hypothetical protein LOC147093 (LOC147093), mRNA
 XM_378676 Homo sapiens LOC400598 (LOC400598), mRNA
 XM_378678 Homo sapiens hypothetical gene supported by AK055254; BC051705 (LOC400599), mRNA
 XM_378680 Homo sapiens hypothetical protein LOC147080 (LOC147080), mRNA
 XM_378682 Homo sapiens LOC400602 (LOC400602), mRNA
 XM_378683 Homo sapiens hypothetical gene supported by AK000454 (LOC400603), mRNA
 XM_378684 Homo sapiens hypothetical gene supported by BC039684 (LOC400604), mRNA
 XM_378685 Homo sapiens hypothetical gene supported by BC039328 (LOC400605), mRNA
 XM_378686 Homo sapiens hypothetical gene supported by AK126827 (LOC400606), mRNA
 XM_378687 Homo sapiens hypothetical protein LOC339210 (LOC339210), mRNA
 XM_378688 Homo sapiens LOC400607 (LOC400607), mRNA
 XM_378689 Homo sapiens LOC400611 (LOC400611), mRNA
 XM_378692 Homo sapiens hypothetical gene supported by AK094767 (LOC400612), mRNA
 XM_378693 Homo sapiens LOC400614 (LOC400614), mRNA
 XM_378694 Homo sapiens hypothetical protein LOC146784 (LOC146784), mRNA
 XM_378695 Homo sapiens hypothetical gene supported by BC053686 (LOC400616), mRNA
 XM_378698 Homo sapiens hypothetical gene supported by AK093963 (LOC400617), mRNA
 XM_378700 Homo sapiens hypothetical gene supported by AK094963 (LOC400618), mRNA
 XM_378701 Homo sapiens hypothetical protein LOC146795 (LOC146795), mRNA
 XM_378703 Homo sapiens hypothetical gene supported by AK129994 (LOC400619), mRNA
 XM_378705 Homo sapiens hypothetical gene supported by BC035399 (LOC400620), mRNA
 XM_378706 Homo sapiens LOC400621 (LOC400621), mRNA
 XM_378708 Homo sapiens hypothetical gene supported by AK130926 (LOC400623), mRNA
 XM_378709 Homo sapiens hypothetical gene supported by AK127023 (LOC400624), mRNA
 XM_378712 Homo sapiens hypothetical protein LOC146713 (LOC146713), mRNA
 XM_378713 Homo sapiens hypothetical gene supported by BX648922 (LOC400626), mRNA
 XM_378723 Homo sapiens hypothetical protein FLJ22659 (FLJ22659), mRNA
 XM_378724 Homo sapiens hypothetical gene supported by AL832615 (LOC400630), mRNA
 XM_378727 Homo sapiens hypothetical gene supported by BC053686 (LOC400632), mRNA
 XM_378728 Homo sapiens LOC400633 (LOC400633), mRNA
 XM_378730 Homo sapiens LOC400638 (LOC400638), mRNA

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XM_378734 Homo sapiens hypothetical protein LOC284214 (LOC284214), mRNA
 XM_378735 Homo sapiens hypothetical gene supported by BC041875; BX648984 (LOC4
 XM_378738 Homo sapiens hypothetical gene supported by AK095347 (LOC400643), mR
 XM_378741 Homo sapiens hypothetical gene supported by AK126243 (LOC400644), mR
 XM_378742 Homo sapiens hypothetical gene supported by AK127888 (LOC400645), mR
 XM_378743 Homo sapiens hypothetical gene supported by BX640930 (LOC400647), mR
 XM_378745 Homo sapiens hypothetical gene supported by BC031271 (LOC400648), mR
 XM_378746 Homo sapiens hypothetical gene supported by AK126075 (LOC400649), mR
 XM_378747 Homo sapiens hypothetical gene supported by AK094936 (LOC100651), mR
 XM_378750 Homo sapiens hypothetical gene supported by BC047606 (LOC400653), mR
 XM_378751 Homo sapiens hypothetical gene supported by BC042493 (LOC400654), mR
 XM_378753 Homo sapiens hypothetical gene supported by BC013370; BC034583 (LOC4
 XM_378754 Homo sapiens hypothetical gene supported by BC039507 (LOC400656), mR
 XM_378755 Homo sapiens hypothetical gene supported by BC036588 (LOC400657), mR
 XM_378756 Homo sapiens hypothetical protein LOC284274 (LOC284274), mRNA
 XM_378757 Homo sapiens hypothetical gene supported by AK093936 (LOC400659), mR
 XM_378758 Homo sapiens hypothetical gene supported by AK094957 (LOC400660), mR
 XM_378760 Homo sapiens hypothetical gene supported by AK055411 (LOC400662), mR
 XM_378763 Homo sapiens hypothetical protein LOC284240 (LOC284240), mRNA
 XM_378765 Homo sapiens hypothetical gene supported by AK096031 (LOC400667), mR
 XM_378766 Homo sapiens LOC400669 (LOC400669), mRNA
 XM_378767 Homo sapiens LOC400670 (LOC400670), mRNA
 XM_378769 Homo sapiens LOC400671 (LOC400671), mRNA
 XM_378770 Homo sapiens hypothetical gene supported by AK001151 (LOC400672), mR
 XM_378776 Homo sapiens LOC400675 (LOC400675), mRNA
 XM_378777 Homo sapiens hypothetical protein LOC284385 (LOC284385), mRNA
 XM_378780 Homo sapiens hypothetical protein LOC126536 (LOC126536), mRNA
 XM_378783 Homo sapiens hypothetical gene supported by AK097381; BC040866 (LOC4
 XM_378784 Homo sapiens hypothetical gene supported by BC030765 (LOC400683), mR
 XM_378786 Homo sapiens hypothetical protein LOC148145 (LOC148145), mRNA
 XM_378787 Homo sapiens hypothetical protein LOC284395 (LOC284395), mRNA
 XM_378791 Homo sapiens hypothetical protein LOC339316 (LOC339316), mRNA
 XM_378793 Homo sapiens hypothetical gene supported by BC000922 (LOC400684), mR
 XM_378794 Homo sapiens hypothetical protein LOC284402 (LOC284402), mRNA
 XM_378795 Homo sapiens hypothetical gene supported by BC045806 (LOC400685), mR
 XM_378796 Homo sapiens hypothetical gene supported by AK125858 (LOC400686), mR
 XM_378798 Homo sapiens hypothetical gene supported by AK092138 (LOC400689), mR
 XM_378799 Homo sapiens LOC400691 (LOC400691), mRNA
 XM_378800 Homo sapiens hypothetical gene supported by BC042546 (LOC400694), mR
 XM_378801 Homo sapiens LOC400695 (LOC400695), mRNA
 XM_378804 Homo sapiens LOC400700 (LOC400700), mRNA
 XM_378805 Homo sapiens LOC400701 (LOC400701), mRNA
 XM_378806 Homo sapiens hypothetical gene supported by AK024119 (LOC400702), mR
 XM_378807 Homo sapiens hypothetical gene supported by AK054869 (LOC400704), mR
 XM_378810 Homo sapiens hypothetical gene supported by AK096622 (LOC400706), mR
 XM_378812 Homo sapiens hypothetical gene supported by AK130360 (LOC400710), mR
 XM_378815 Homo sapiens LOC400722 (LOC400722), mRNA
 XM_378820 Homo sapiens hypothetical gene supported by AK097327; BC037297 (LOC4
 XM_378822 Homo sapiens hypothetical protein LOC254099 (LOC254099), mRNA
 XM_378823 Homo sapiens hypothetical protein LOC148413 (LOC148413), mRNA
 XM_378824 Homo sapiens LOC400729 (LOC400729), mRNA
 XM_378825 Homo sapiens hypothetical gene supported by AK097814 (LOC400730), mR
 XM_378828 Homo sapiens hypothetical protein LOC115110 (LOC115110), mRNA
 XM_378831 Homo sapiens hypothetical gene supported by AK124708 (LOC400732), mR
 XM_378832 Homo sapiens hypothetical protein LOC284661 (LOC284661), mRNA
 XM_378835 Homo sapiens LOC400733 (LOC400733), mRNA
 XM_378837 Homo sapiens hypothetical gene supported by AK091499 (LOC400738), mR

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XM_378838 Homo sapiens hypothetical gene supported by AK125737 (LOC400739), mRNA
 XM_378840 Homo sapiens hypothetical gene supported by BC036435 (LOC400740), mRNA
 XM_378841 Homo sapiens LOC400741 (LOC400741), mRNA
 XM_378842 Homo sapiens hypothetical gene supported by BC033316 (LOC400742), mRNA
 XM_378843 Homo sapiens hypothetical gene supported by AK127830 (LOC400743), mRNA
 XM_378846 Homo sapiens PNAS-123 (LOC85028), mRNA
 XM_378852 Homo sapiens hypothetical gene supported by BC040627 (LOC400748), mRNA
 XM_378855 Homo sapiens hypothetical protein LOC339442 (LOC339442), mRNA
 XM_378858 Homo sapiens hypothetical protein LOC339539 (LOC339539), mRNA
 XM_378859 Homo sapiens hypothetical gene supported by BC031250 (LOC400751), mRNA
 XM_378860 Homo sapiens hypothetical protein LOC149478 (LOC149478), mRNA
 XM_378861 Homo sapiens hypothetical gene supported by BC006119 (LOC400752), mRNA
 XM_378862 Homo sapiens LOC400753 (LOC400753), mRNA
 XM_378865 Homo sapiens hypothetical gene supported by BC030752 (LOC400756), mRNA
 XM_378866 Homo sapiens hypothetical protein LOC199899 (LOC199899), mRNA
 XM_378873 Homo sapiens LOC400758 (LOC400758), mRNA
 XM_378874 Homo sapiens hypothetical gene supported by AK130864 (LOC400761), mRNA
 XM_378876 Homo sapiens hypothetical protein LOC149351 (LOC149351), mRNA
 XM_378877 Homo sapiens hypothetical gene supported by AL832786 (LOC400762), mRNA
 XM_378879 Homo sapiens hypothetical gene supported by AK000394 (LOC400763), mRNA
 XM_378880 Homo sapiens hypothetical gene supported by AK094798 (LOC400764), mRNA
 XM_378883 Homo sapiens hypothetical gene supported by BC051808 (LOC400768), mRNA
 XM_378886 Homo sapiens hypothetical protein LOC284475 (LOC284475), mRNA
 XM_378889 Homo sapiens hypothetical gene supported by AK090412 (LOC400770), mRNA
 XM_378890 Homo sapiens hypothetical gene supported by AK098337; BC022881 (LOC400771), mRNA
 XM_378891 Homo sapiens hypothetical gene supported by BC012753 (LOC400772), mRNA
 XM_378892 Homo sapiens LOC400775 (LOC400775), mRNA
 XM_378893 Homo sapiens hypothetical gene supported by AK125616 (LOC400777), mRNA
 XM_378894 Homo sapiens LOC400778 (LOC400778), mRNA
 XM_378897 Homo sapiens LOC400781 (LOC400781), mRNA
 XM_378898 Homo sapiens LOC400782 (LOC400782), mRNA
 XM_378899 Homo sapiens LOC400783 (LOC400783), mRNA
 XM_378901 Homo sapiens hypothetical gene supported by M60502 (LOC400786), mRNA
 XM_378903 Homo sapiens LOC400789 (LOC400789), mRNA
 XM_378905 Homo sapiens hypothetical gene supported by AK094742; AK128347 (LOC400790), mRNA
 XM_378908 Homo sapiens hypothetical gene supported by AK125122 (LOC400793), mRNA
 XM_378909 Homo sapiens hypothetical gene supported by BC030596 (LOC400794), mRNA
 XM_378910 Homo sapiens hypothetical gene supported by AK000073 (LOC400795), mRNA
 XM_378912 Homo sapiens hypothetical protein LOC284688 (LOC284688), mRNA
 XM_378914 Homo sapiens KIAA0492 protein (KIAA0492), mRNA
 XM_378917 Homo sapiens LOC400796 (LOC400796), mRNA
 XM_378918 Homo sapiens hypothetical gene supported by AK125993 (LOC400797), mRNA
 XM_378919 Homo sapiens hypothetical gene supported by AK092849 (LOC400798), mRNA
 XM_378921 Homo sapiens hypothetical protein LOC148756 (LOC148756), mRNA
 XM_378923 Homo sapiens hypothetical protein LOC339476 (LOC339476), mRNA
 XM_378925 Homo sapiens hypothetical gene supported by AK125573 (LOC400800), mRNA
 XM_378930 Homo sapiens hypothetical gene supported by AK097184 (LOC400801), mRNA
 XM_378933 Homo sapiens LOC400802 (LOC400802), mRNA
 XM_378934 Homo sapiens LOC400803 (LOC400803), mRNA
 XM_378941 Homo sapiens hypothetical protein LOC339535 (LOC339535), mRNA
 XM_378945 Homo sapiens hypothetical protein LOC149134 (LOC149134), mRNA
 XM_378946 Homo sapiens hypothetical gene supported by AK096414 (LOC400812), mRNA
 XM_378947 Homo sapiens LOC400813 (LOC400813), mRNA
 XM_378949 Homo sapiens similar to hypothetical protein LOC148413 (LOC400817), mRNA
 XM_378950 Homo sapiens hypothetical gene supported by AK097614 (LOC400819), mRNA
 XM_378951 Homo sapiens hypothetical protein LOC284628 (LOC284628), mRNA
 XM_378954 Homo sapiens hypothetical gene supported by AK125616 (LOC400828), mRNA

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XM_378955 Homo sapiens LOC400829 (LOC400829), mRNA
 XM_378956 Homo sapiens hypothetical gene supported by AK092524 (LOC400831), mR
 XM_378957 Homo sapiens LOC400832 (LOC400832), mRNA
 XM_378961 Homo sapiens LOC400835 (LOC400835), mRNA
 XM_378964 Homo sapiens hypothetical protein LOC339593 (LOC339593), mRNA
 XM_378969 Homo sapiens hypothetical protein LOC284788 (LOC284788), mRNA
 XM_378970 Homo sapiens hypothetical gene supported by AK090900 (LOC400839), mR
 XM_378971 Homo sapiens hypothetical protein LOC284798 (LOC284798), mRNA
 XM_378973 Homo sapiens hypothetical protein LOC284901 (LOC284901), mRNA
 XM_378974 Homo sapiens hypothetical gene supported by AK127732 (LOC400841), mR
 XM_378976 Homo sapiens LOC400843 (LOC400843), mRNA
 XM_378977 Homo sapiens hypothetical gene supported by AK124127 (LOC400844), mR
 XM_378978 Homo sapiens hypothetical gene supported by AK090932 (LOC400845), mR
 XM_378980 Homo sapiens hypothetical protein LOC339568 (LOC339568), mRNA
 XM_378981 Homo sapiens hypothetical gene supported by AK127953 (LOC400846), mR
 XM_378982 Homo sapiens hypothetical gene supported by AL832259; BC035184 (LOC4
 XM_378983 Homo sapiens hypothetical protein LOC284749 (LOC284749), mRNA
 XM_378985 Homo sapiens hypothetical protein LOC284751 (LOC284751), mRNA
 XM_378988 Homo sapiens LOC400849 (LOC400849), mRNA
 XM_378989 Homo sapiens hypothetical gene supported by AK126744 (LOC400850), mR
 XM_378992 Homo sapiens hypothetical gene supported by BC015673 (LOC400851), mR
 XM_378993 Homo sapiens hypothetical gene supported by AK124784 (LOC400852), mR
 XM_378994 Homo sapiens hypothetical gene supported by AK125376 (LOC400853), mR
 XM_379002 Homo sapiens LOC400861 (LOC400861), mRNA
 XM_379003 Homo sapiens LOC400862 (LOC400862), mRNA
 XM_379004 Homo sapiens hypothetical gene supported by AK127913 (LOC400863), mR
 XM_379006 Homo sapiens hypothetical gene supported by AK094771 (LOC400865), mR
 XM_379009 Homo sapiens hypothetical gene supported by AY204750 (LOC400866), mR
 XM_379011 Homo sapiens hypothetical protein LOC284835 (LOC284835), mRNA
 XM_379012 Homo sapiens hypothetical gene supported by BC033280 (LOC400868), mR
 XM_379014 Homo sapiens LOC400869 (LOC400869), mRNA
 XM_379015 Homo sapiens hypothetical gene supported by AK123727 (LOC400870), mR
 XM_379016 Homo sapiens hypothetical gene supported by AK057962 (LOC400871), mR
 XM_379017 Homo sapiens LOC400872 (LOC400872), mRNA
 XM_379018 Homo sapiens hypothetical gene supported by BX648824 (LOC400873), mR
 XM_379020 Homo sapiens hypothetical gene supported by AK096288 (LOC400874), mR
 XM_379021 Homo sapiens chromosome 21 open reading frame 30 (C2orf30), mRNA
 XM_379022 Homo sapiens hypothetical gene supported by BC040064 (LOC400875), mR
 XM_379023 Homo sapiens hypothetical gene supported by BC036902 (LOC400876), mR
 XM_379025 Homo sapiens LOC400877 (LOC400877), mRNA
 XM_379029 Homo sapiens hypothetical gene supported by AK099951 (LOC400879), mR
 XM_379030 Homo sapiens hypothetical gene supported by AY338954 (LOC400880), mR
 XM_379032 Homo sapiens hypothetical gene supported by BC021738 (LOC400885), mR
 XM_379036 Homo sapiens LOC400890 (LOC400890), mRNA
 XM_379040 Homo sapiens hypothetical gene supported by AK097628 (LOC400919), mR
 XM_379041 Homo sapiens hypothetical gene supported by BC040576 (LOC400920), mR
 XM_379044 Homo sapiens hypothetical protein LOC284898 (LOC284898), mRNA
 XM_379046 Homo sapiens LOC400923 (LOC400923), mRNA
 XM_379052 Homo sapiens hypothetical gene supported by AK075161 (LOC400926), mR
 XM_379054 Homo sapiens hypothetical protein LOC339674 (LOC339674), mRNA
 XM_379055 Homo sapiens hypothetical gene supported by AK130208 (LOC400929), mR
 XM_379060 Homo sapiens hypothetical protein LOC339685 (LOC339685), mRNA
 XM_379064 Homo sapiens hypothetical protein LOC284933 (LOC284933), mRNA
 XM_379065 Homo sapiens hypothetical gene supported by BC033837 (LOC400933), mR
 XM_379068 Homo sapiens hypothetical gene supported by BC055007 (LOC400937), mR
 XM_379069 Homo sapiens hypothetical protein LOC339822 (LOC339822), mRNA
 XM_379072 Homo sapiens LOC400939 (LOC400939), mRNA

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XM_379073 Homo sapiens hypothetical protein LOC386597 (LOC386597), mRNA
 XM_379074 Homo sapiens hypothetical protein LOC339788 (LOC339788), mRNA
 XM_379075 Homo sapiens hypothetical gene supported by BX641130 (LOC400942), mR
 XM_379077 Homo sapiens LOC400944 (LOC400944), mRNA
 XM_379078 Homo sapiens hypothetical gene supported by AK123475 (LOC400945), mR
 XM_379079 Homo sapiens hypothetical gene supported by AK022396; AK097927 (LOC4
 XM_379080 Homo sapiens LOC400947 (LOC400947), mRNA
 XM_379085 Homo sapiens hypothetical protein LOC285043 (LOC285043), mRNA
 XM_379086 Homo sapiens hypothetical protein LOC285045 (LOC285045), mRNA
 XM_379089 Homo sapiens LOC400951 (LOC400951), mRNA
 XM_379094 Homo sapiens hypothetical gene supported by BX647332 (LOC400953), mR
 XM_379096 Homo sapiens hypothetical gene supported by AL832565 (LOC400955), mR
 XM_379097 Homo sapiens hypothetical gene supported by AK122786 (LOC400957), mR
 XM_379098 Homo sapiens hypothetical protein LOC339803 (LOC339803), mRNA
 XM_379099 Homo sapiens hypothetical protein LOC339807 (LOC339807), mRNA
 XM_379100 Homo sapiens hypothetical gene supported by BC037562 (LOC400958), mR
 XM_379101 Homo sapiens hypothetical gene supported by BC033059 (LOC400959), mR
 XM_379102 Homo sapiens hypothetical gene supported by BC040598 (LOC400960), mR
 XM_379106 Homo sapiens hypothetical gene supported by BC044795 (LOC400964), mR
 XM_379108 Homo sapiens LOC400969 (LOC400969), mRNA
 XM_379109 Homo sapiens hypothetical gene supported by AK127783 (LOC400983), mR
 XM_379111 Homo sapiens hypothetical protein LOC285033 (LOC285033), mRNA
 XM_379112 Homo sapiens LOC400988 (LOC400988), mRNA
 XM_379113 Homo sapiens hypothetical gene supported by BC040181 (LOC400990), mR
 XM_379114 Homo sapiens hypothetical protein LOC150577 (LOC150577), mRNA
 XM_379117 Homo sapiens hypothetical protein LOC150568 (LOC150568), mRNA
 XM_379118 Homo sapiens hypothetical gene supported by AK095498 (LOC400992), mR
 XM_379119 Homo sapiens hypothetical protein LOC285000 (LOC285000), mRNA
 XM_379121 Homo sapiens hypothetical gene supported by AK056084; AK095678; NM_1
 XM_379122 Homo sapiens hypothetical gene supported by AK125994 (LOC400997), mR
 XM_379123 Homo sapiens hypothetical gene supported by AK124342 (LOC400999), mR
 XM_379131 Homo sapiens LOC401001 (LOC401001), mRNA
 XM_379133 Homo sapiens hypothetical protein LOC151121 (LOC151121), mRNA
 XM_379135 Homo sapiens hypothetical gene supported by AK093281 (LOC401005), mR
 XM_379136 Homo sapiens hypothetical gene supported by AK093281 (LOC401006), mR
 XM_379141 Homo sapiens hypothetical gene supported by BC043549; BX648102 (LOC4
 XM_379145 Homo sapiens hypothetical gene supported by AK092134 (LOC401020), mR
 XM_379146 Homo sapiens hypothetical gene supported by BC047605 (LOC401021), mR
 XM_379147 Homo sapiens hypothetical gene supported by BC030713; BC047481 (LOC4
 XM_379149 Homo sapiens LOC401025 (LOC401025), mRNA
 XM_379154 Homo sapiens hypothetical protein LOC151300 (LOC151300), mRNA
 XM_379156 Homo sapiens LOC401032 (LOC401032), mRNA
 XM_379158 Homo sapiens hypothetical gene supported by AK055016 (LOC401033), mR
 XM_379159 Homo sapiens hypothetical protein LOC151484 (LOC151484), mRNA
 XM_379161 Homo sapiens hypothetical gene supported by AK056246 (LOC401037), mR
 XM_379163 Homo sapiens hypothetical gene supported by AK057585 (LOC401038), mR
 XM_379164 Homo sapiens hypothetical protein LOC151171 (LOC151171), mRNA
 XM_379166 Homo sapiens hypothetical gene supported by AK056439 (LOC401041), mR
 XM_379167 Homo sapiens LOC401042 (LOC401042), mRNA
 XM_379168 Homo sapiens LOC401043 (LOC401043), mRNA
 XM_379169 Homo sapiens hypothetical gene supported by AK098031 (LOC401044), mR
 XM_379170 Homo sapiens LOC401046 (LOC401046), mRNA
 XM_379171 Homo sapiens LOC401048 (LOC401048), mRNA
 XM_379172 Homo sapiens hypothetical gene supported by AK124857 (LOC401049), mR
 XM_379173 Homo sapiens hypothetical gene supported by AK124088 (LOC401050), mR
 XM_379174 Homo sapiens LOC401051 (LOC401051), mRNA
 XM_379175 Homo sapiens hypothetical gene supported by AK022260 (LOC401052), mR

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XM_379177 Homo sapiens hypothetical gene supported by BC041425 (LOC401053), mR
XM_379179 Homo sapiens hypothetical protein LOC152274 (LOC152274), mRNA
XM_379180 Homo sapiens LOC401056 (LOC401056), mRNA
XM_379181 Homo sapiens LOC401057 (LOC401057), mRNA
XM_379182 Homo sapiens LOC401058 (LOC401058), mRNA
XM_379183 Homo sapiens hypothetical protein LOC152024 (LOC152024), mRNA
XM_379184 Homo sapiens hypothetical gene supported by AK096885; AK098084 (LOC4
XM_379189 Homo sapiens hypothetical gene supported by AK057338 (LOC401061), mR
XM_379190 Homo sapiens hypothetical gene supported by AK092973 (LOC401062), mR
XM_379191 Homo sapiens LOC401065 (LOC401065), mRNA
XM_379192 Homo sapiens LOC401066 (LOC401066), mRNA
XM_379194 Homo sapiens hypothetical gene supported by BC028186 (LOC401068), mR
XM_379195 Homo sapiens hypothetical protein LOC285401 (LOC285401), mRNA
XM_379196 Homo sapiens hypothetical protein LOC151877 (LOC151877), mRNA
XM_379197 Homo sapiens hypothetical gene supported by AL832401 (LOC401073), mR
XM_379198 Homo sapiens hypothetical protein LOC285286 (LOC285286), mRNA
XM_379200 Homo sapiens LOC401078 (LOC401078), mRNA
XM_379201 Homo sapiens LOC401079 (LOC401079), mRNA
XM_379203 Homo sapiens hypothetical protein LOC348801 (LOC348801), mRNA
XM_379204 Homo sapiens hypothetical protein LOC152225 (LOC152225), mRNA
XM_379205 Homo sapiens hypothetical protein LOC151658 (LOC151658), mRNA
XM_379206 Homo sapiens hypothetical gene supported by AK026416 (LOC401081), mR
XM_379207 Homo sapiens hypothetical protein LOC285194 (LOC285194), mRNA
XM_379210 Homo sapiens LOC401085 (LOC401085), mRNA
XM_379213 Homo sapiens LOC401086 (LOC401086), mRNA
XM_379214 Homo sapiens hypothetical protein LOC339942 (LOC339942), mRNA
XM_379215 Homo sapiens hypothetical protein LOC132241 (LOC132241), mRNA
XM_379228 Homo sapiens LOC401093 (LOC401093), mRNA
XM_379229 Homo sapiens LOC401094 (LOC401094), mRNA
XM_379230 Homo sapiens hypothetical protein LOC339894 (LOC339894), mRNA
XM_379231 Homo sapiens hypothetical gene supported by BC034803 (LOC401098), mR
XM_379233 Homo sapiens LOC401099 (LOC401099), mRNA
XM_379234 Homo sapiens LOC401101 (LOC401101), mRNA
XM_379235 Homo sapiens hypothetical gene supported by AK127955 (LOC401103), mR
XM_379240 Homo sapiens LOC401104 (LOC401104), mRNA
XM_379243 Homo sapiens hypothetical gene supported by AK091527 (LOC401106), mR
XM_379244 Homo sapiens hypothetical gene supported by AK127609 (LOC401107), mR
XM_379247 Homo sapiens hypothetical gene supported by AK128780; AL137733 (LOC4
XM_379248 Homo sapiens hypothetical gene supported by AK098259 (LOC401112), mR
XM_379249 Homo sapiens hypothetical gene supported by AK123125 (LOC401113), mR
XM_379250 Homo sapiens hypothetical gene supported by BC038466; BC062790 (LOC4
XM_379252 Homo sapiens hypothetical gene supported by BC017173 (LOC401117), mR
XM_379254 Homo sapiens hypothetical protein LOC339988 (LOC339988), mRNA
XM_379255 Homo sapiens hypothetical gene supported by AK094096 (LOC401119), mR
XM_379256 Homo sapiens hypothetical gene supported by AK127863 (LOC401120), mR
XM_379258 Homo sapiens hypothetical protein LOC285547 (LOC285547), mRNA
XM_379260 Homo sapiens hypothetical protein LOC152742 (LOC152742), mRNA
XM_379262 Homo sapiens LOC401124 (LOC401124), mRNA
XM_379263 Homo sapiens LOC401126 (LOC401126), mRNA
XM_379264 Homo sapiens LOC401128 (LOC401128), mRNA
XM_379265 Homo sapiens LOC401129 (LOC401129), mRNA
XM_379267 Homo sapiens hypothetical gene supported by BC040544 (LOC401134), mR
XM_379268 Homo sapiens hypothetical gene supported by AK093682; AK129519 (LOC4
XM_379270 Homo sapiens caseln alpha s2-like A (CSN1S2A), mRNA
XM_379271 Homo sapiens LOC401144 (LOC401144), mRNA
XM_379273 Homo sapiens hypothetical gene supported by AK024248; AL137733 (LOC4
XM_379274 Homo sapiens hypothetical gene supported by BC062741 (LOC401151), mR

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XM_379275 Homo sapiens LOC401153 (LOC401153), mRNA
 XM_379276 Homo sapiens hypothetical gene supported by AK123449; BX84 1014 (LOC4
 XM_379278 Homo sapiens LOC401156 (LOC401156), mRNA
 XM_379280 Homo sapiens hypothetical protein LOC285422 (LOC285422), mRNA
 XM_379287 Homo sapiens LOC401159 (LOC401159), mRNA
 XM_379288 Homo sapiens hypothetical protein LOC340017 (LOC340017), mRNA
 XM_379294 Homo sapiens similar to hypothetical protein DKFp762K222 (LOC401163),
 XM_379295 Homo sapiens hypothetical protein LOC285441 (LOC285441), mRNA
 XM_379298 Homo sapiens hypothetical gene supported by AY494056 (LOC401164), mR
 XM_379299 Homo sapiens hypothetical gene supported by BC029568 (LOC401165), mR
 XM_379303 Homo sapiens hypothetical gene supported by AK126199 (LOC401168), mR
 XM_379306 Homo sapiens hypothetical gene supported by BC034612 (LOC401169), mR
 XM_379309 Homo sapiens hypothetical gene supported by BC035019 (LOC401171), mR
 XM_379313 Homo sapiens hypothetical gene supported by AK126802 (LOC401173), mR
 XM_379317 Homo sapiens LOC401175 (LOC401175), mRNA
 XM_379318 Homo sapiens hypothetical gene supported by BC043001 (LOC401176), mR
 XM_379320 Homo sapiens hypothetical gene supported by BC052942 (LOC401177), mR
 XM_379321 Homo sapiens hypothetical protein LOC340107 (LOC340107), mRNA
 XM_379322 Homo sapiens hypothetical protein LOC340109 (LOC340109), mRNA
 XM_379323 Homo sapiens LOC401178 (LOC401178), mRNA
 XM_379324 Homo sapiens hypothetical protein LOC340113 (LOC340113), mRNA
 XM_379325 Homo sapiens LOC401180 (LOC401180), mRNA
 XM_379326 Homo sapiens hypothetical gene supported by BC028978 (LOC401181), mR
 XM_379327 Homo sapiens LOC401182 (LOC401182), mRNA
 XM_379328 Homo sapiens LOC401184 (LOC401184), mRNA
 XM_379331 Homo sapiens hypothetical gene supported by AK057759 (LOC401185), mR
 XM_379332 Homo sapiens LOC401186 (LOC401186), mRNA
 XM_379333 Homo sapiens LOC401187 (LOC401187), mRNA
 XM_379334 Homo sapiens hypothetical protein LOC257396 (LOC257396), mRNA
 XM_379336 Homo sapiens hypothetical gene supported by AK091013 (LOC401188), mR
 XM_379339 Homo sapiens LOC401199 (LOC401199), mRNA
 XM_379340 Homo sapiens hypothetical protein LOC285713 (LOC285713), mRNA
 XM_379343 Homo sapiens hypothetical gene supported by AK092258 (LOC401201), mR
 XM_379347 Homo sapiens LOC401203 (LOC401203), mRNA
 XM_379355 Homo sapiens hypothetical protein LOC340074 (LOC340074), mRNA
 XM_379359 Homo sapiens hypothetical gene supported by AK022326 (LOC401210), mR
 XM_379363 Homo sapiens hypothetical gene supported by BX840700 (LOC401212), mR
 XM_379364 Homo sapiens hypothetical gene supported by AK127910 (LOC401213), mR
 XM_379366 Homo sapiens hypothetical gene supported by AK092848; AK123816 (LOC4
 XM_379368 Homo sapiens LOC401215 (LOC401215), mRNA
 XM_379371 Homo sapiens hypothetical protein LOC285626 (LOC285626), mRNA
 XM_379372 Homo sapiens hypothetical gene supported by BC039501 (LOC401216), mR
 XM_379373 Homo sapiens hypothetical protein LOC257358 (LOC257358), mRNA
 XM_379377 Homo sapiens hypothetical gene supported by BX849016 (LOC401219), mR
 XM_379378 Homo sapiens hypothetical gene supported by BC036933 (LOC401220), mR
 XM_379380 Homo sapiens hypothetical gene supported by BC036933 (LOC401222), mR
 XM_379381 Homo sapiens hypothetical gene supported by AK055745 (LOC401225), mR
 XM_379382 Homo sapiens hypothetical gene supported by AK093197; BC040992 (LOC4
 XM_379384 Homo sapiens hypothetical protein LOC285766 (LOC285766), mRNA
 XM_379386 Homo sapiens hypothetical protein LOC285768 (LOC285768), mRNA
 XM_379391 Homo sapiens hypothetical gene supported by AK128409 (LOC401230), mR
 XM_379392 Homo sapiens hypothetical gene supported by AK023629 (LOC401231), mR
 XM_379393 Homo sapiens hypothetical gene supported by BX840708 (LOC401232), mR
 XM_379395 Homo sapiens hypothetical gene supported by BC014487 (LOC401234), mR
 XM_379396 Homo sapiens hypothetical protein LOC221710 (LOC221710), mRNA
 XM_379398 Homo sapiens hypothetical gene supported by AK028189 (LOC401237), mR
 XM_379401 Homo sapiens LOC401241 (LOC401241), mRNA

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XM_379402 Homo sapiens hypothetical gene supported by AK055503 (LOC401242), mRNA
 XM_379403 Homo sapiens chromosome 6 open reading frame 12 (C6orf12), mRNA
 XM_379404 Homo sapiens hypothetical gene supported by AK055657 (LOC401245), mRNA
 XM_379406 Homo sapiens hypothetical gene supported by AL832447 (LOC401254), mRNA
 XM_379408 Homo sapiens LOC401255 (LOC401255), mRNA
 XM_379409 Homo sapiens hypothetical gene supported by BX648112 (LOC401256), mRNA
 XM_379410 Homo sapiens hypothetical gene supported by BC034770 (LOC401257), mRNA
 XM_379411 Homo sapiens hypothetical gene supported by AK125083 (LOC401258), mRNA
 XM_379413 Homo sapiens hypothetical gene supported by AK098234 (LOC401261), mRNA
 XM_379417 Homo sapiens hypothetical gene supported by AK095117 (LOC401264), mRNA
 XM_379424 Homo sapiens LOC401268 (LOC401268), mRNA
 XM_379430 Homo sapiens hypothetical protein LOC285758 (LOC285758), mRNA
 XM_379432 Homo sapiens hypothetical protein LOC285733 (LOC285733), mRNA
 XM_379433 Homo sapiens hypothetical protein LOC285735 (LOC285735), mRNA
 XM_379434 Homo sapiens hypothetical protein LOC154092 (LOC154092), mRNA
 XM_379435 Homo sapiens hypothetical gene supported by AK128874 (LOC401275), mRNA
 XM_379436 Homo sapiens hypothetical gene supported by BC038188 (LOC401276), mRNA
 XM_379437 Homo sapiens hypothetical protein LOC153910 (LOC153910), mRNA
 XM_379438 Homo sapiens hypothetical protein LOC285740 (LOC285740), mRNA
 XM_379439 Homo sapiens LOC401277 (LOC401277), mRNA
 XM_379441 Homo sapiens LOC401279 (LOC401279), mRNA
 XM_379450 Homo sapiens hypothetical gene supported by AK130765 (LOC401281), mRNA
 XM_379452 Homo sapiens hypothetical gene supported by AL831931 (LOC401282), mRNA
 XM_379453 Homo sapiens hypothetical gene supported by AL832143 (LOC401283), mRNA
 XM_379454 Homo sapiens hypothetical gene supported by AK095077 (LOC401284), mRNA
 XM_379456 Homo sapiens hypothetical protein LOC154222 (LOC154222), mRNA
 XM_379458 Homo sapiens hypothetical gene supported by BX648586 (LOC401287), mRNA
 XM_379459 Homo sapiens hypothetical gene supported by AK095441 (LOC401289), mRNA
 XM_379460 Homo sapiens hypothetical gene supported by AL832760 (LOC401290), mRNA
 XM_379461 Homo sapiens LOC401291 (LOC401291), mRNA
 XM_379462 Homo sapiens hypothetical gene supported by BC084362; BX537893 (LOC401292), mRNA
 XM_379463 Homo sapiens similar to hypothetical protein LOC154222 (LOC401294), mRNA
 XM_379467 Homo sapiens hypothetical gene supported by AK125766 (LOC401296), mRNA
 XM_379469 Homo sapiens hypothetical gene supported by BC032734 (LOC401297), mRNA
 XM_379471 Homo sapiens LOC401299 (LOC401299), mRNA
 XM_379472 Homo sapiens LOC401301 (LOC401301), mRNA
 XM_379474 Homo sapiens hypothetical gene supported by AK127500 (LOC401310), mRNA
 XM_379476 Homo sapiens hypothetical gene supported by BC039682 (LOC401312), mRNA
 XM_379477 Homo sapiens hypothetical protein LOC285941 (LOC285941), mRNA
 XM_379478 Homo sapiens hypothetical gene supported by AK093987 (LOC401315), mRNA
 XM_379479 Homo sapiens LOC401317 (LOC401317), mRNA
 XM_379480 Homo sapiens LOC401318 (LOC401318), mRNA
 XM_379481 Homo sapiens hypothetical gene supported by BC023581; BC044638 (LOC401319), mRNA
 XM_379482 Homo sapiens hypothetical gene supported by BC016976 (LOC401320), mRNA
 XM_379483 Homo sapiens hypothetical gene supported by AK092714 (LOC401321), mRNA
 XM_379484 Homo sapiens hypothetical gene supported by AL832092 (LOC401324), mRNA
 XM_379485 Homo sapiens LOC401326 (LOC401326), mRNA
 XM_379486 Homo sapiens hypothetical protein LOC285958 (LOC285958), mRNA
 XM_379487 Homo sapiens hypothetical gene supported by AK125311 (LOC401334), mRNA
 XM_379488 Homo sapiens hypothetical gene supported by AK024248; AL137733; BC061248 (LOC401335), mRNA
 XM_379489 Homo sapiens LOC401345 (LOC401345), mRNA
 XM_379490 Homo sapiens LOC401346 (LOC401346), mRNA
 XM_379491 Homo sapiens LOC401348 (LOC401348), mRNA
 XM_379492 Homo sapiens hypothetical gene supported by BX648489 (LOC401349), mRNA
 XM_379493 Homo sapiens LOC401352 (LOC401352), mRNA
 XM_379494 Homo sapiens LOC401353 (LOC401353), mRNA
 XM_379495 Homo sapiens LOC401358 (LOC401358), mRNA

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XM_379496 Homo sapiens LOC401359 (LOC401359), mRNA
 XM_379498 Homo sapiens LOC401363 (LOC401363), mRNA
 XM_379499 Homo sapiens hypothetical gene supported by AK024371; BC037920 (LOC4
 XM_379500 Homo sapiens hypothetical gene supported by AK054923; AK126730; NM_0
 XM_379501 Homo sapiens LOC401367 (LOC401367), mRNA
 XM_379502 Homo sapiens LOC401368 (LOC401368), mRNA
 XM_379503 Homo sapiens LOC401371 (LOC401371), mRNA
 XM_379504 Homo sapiens hypothetical gene supported by AK024602 (LOC401380), mR
 XM_379506 Homo sapiens LOC401384 (LOC401384), mRNA
 XM_379507 Homo sapiens LOC401385 (LOC401385), mRNA
 XM_379508 Homo sapiens LOC401386 (LOC401386), mRNA
 XM_379510 Homo sapiens hypothetical protein FLJ34048 (FLJ34048), mRNA
 XM_379511 Homo sapiens LOC401390 (LOC401390), mRNA
 XM_379512 Homo sapiens LOC401394 (LOC401394), mRNA
 XM_379513 Homo sapiens LOC401396 (LOC401396), mRNA
 XM_379514 Homo sapiens hypothetical protein LOC340340 (LOC340340), mRNA
 XM_379515 Homo sapiens hypothetical gene supported by BX537645 (LOC401397), mR
 XM_379516 Homo sapiens hypothetical gene supported by BX648695 (LOC401398), mR
 XM_379517 Homo sapiens hypothetical gene supported by BC063892 (LOC401399), mR
 XM_379518 Homo sapiens LOC401400 (LOC401400), mRNA
 XM_379520 Homo sapiens hypothetical protein FLJ43663 (FLJ43663), mRNA
 XM_379521 Homo sapiens LOC401405 (LOC401405), mRNA
 XM_379522 Homo sapiens LOC401405 (LOC401406), mRNA
 XM_379523 Homo sapiens LOC401407 (LOC401407), mRNA
 XM_379524 Homo sapiens LOC401408 (LOC401408), mRNA
 XM_379526 Homo sapiens hypothetical gene supported by BX648692 (LOC401410), mR
 XM_379527 Homo sapiens hypothetical gene supported by BC023225 (LOC401431), mR
 XM_379528 Homo sapiens hypothetical protein LOC90520 (LOC90520), mRNA
 XM_379529 Homo sapiens LOC401432 (LOC401432), mRNA
 XM_379530 Homo sapiens hypothetical protein LOC285972 (LOC285972), mRNA
 XM_379531 Homo sapiens LOC401434 (LOC401434), mRNA
 XM_379532 Homo sapiens LOC401435 (LOC401435), mRNA
 XM_379533 Homo sapiens LOC401436 (LOC401436), mRNA
 XM_379534 Homo sapiens hypothetical gene supported by AK054822 (LOC401437), mR
 XM_379535 Homo sapiens hypothetical protein LOC285889 (LOC285889), mRNA
 XM_379536 Homo sapiens hypothetical gene supported by BC041429 (LOC401438), mR
 XM_379537 Homo sapiens hypothetical gene supported by AY166699 (LOC401439), mR
 XM_379539 Homo sapiens LOC401440 (LOC401440), mRNA
 XM_379540 Homo sapiens hypothetical protein LOC157693 (LOC157693), mRNA
 XM_379541 Homo sapiens hypothetical gene supported by AK127852 (LOC401441), mR
 XM_379543 Homo sapiens hypothetical gene supported by BC028401 (LOC401442), mR
 XM_379545 Homo sapiens hypothetical gene supported by BC030648 (LOC401443), mR
 XM_379547 Homo sapiens hypothetical gene supported by AK057888 (LOC401445), mR
 XM_379548 Homo sapiens hypothetical gene supported by AK124896; BC037255 (LOC4
 XM_379550 Homo sapiens hypothetical gene supported by AK091259 (LOC401448), mR
 XM_379551 Homo sapiens hypothetical protein LOC349196 (LOC349196), mRNA
 XM_379552 Homo sapiens LOC401449 (LOC401449), mRNA
 XM_379553 Homo sapiens similar to hypothetical protein LOC157278 (LOC401451), mR
 XM_379554 Homo sapiens hypothetical protein LOC157273 (LOC157273), mRNA
 XM_379559 Homo sapiens hypothetical protein LOC157278 (LOC157278), mRNA
 XM_379562 Homo sapiens LOC401456 (LOC401456), mRNA
 XM_379573 Homo sapiens hypothetical protein LOC286135 (LOC286135), mRNA
 XM_379582 Homo sapiens hypothetical protein LOC286177 (LOC286177), mRNA
 XM_379583 Homo sapiens hypothetical gene supported by AK124256 (LOC401462), mR
 XM_379584 Homo sapiens hypothetical gene supported by BC022555; BC050012 (LOC4
 XM_379586 Homo sapiens hypothetical protein LOC286186 (LOC286186), mRNA
 XM_379587 Homo sapiens LOC401464 (LOC401464), mRNA

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XM_379592 Homo sapiens hypothetical protein LOC286144 (LOC286144), mRNA
 XM_379594 Homo sapiens hypothetical protein LOC286149 (LOC286149), mRNA
 XM_379595 Homo sapiens hypothetical gene supported by AK125891 (LOC401471), mR
 XM_379596 Homo sapiens LOC401473 (LOC401473), mRNA
 XM_379597 Homo sapiens hypothetical protein FLJ10489 (FLJ10489), mRNA
 XM_379601 Homo sapiens hypothetical gene supported by BC036187; NM_005839 (LOC
 XM_379603 Homo sapiens hypothetical gene supported by BC009730; BC015157 (LOC4
 XM_379605 Homo sapiens LOC401477 (LOC401477), mRNA
 XM_379608 Homo sapiens hypothetical gene supported by AK056998 (LOC401480), mR
 XM_379609 Homo sapiens hypothetical gene supported by BC041936 (LOC401481), mR
 XM_379610 Homo sapiens LOC401482 (LOC401482), mRNA
 XM_379617 Homo sapiens hypothetical gene supported by AK093004 (LOC401488), mR
 XM_379618 Homo sapiens hypothetical gene supported by AY343891; AY343892; AY34:
 XM_379619 Homo sapiens hypothetical gene supported by BC052949 (LOC401490), mR
 XM_379622 Homo sapiens hypothetical gene supported by AK092343 (LOC401491), mR
 XM_379623 Homo sapiens hypothetical gene supported by AK123194 (LOC401492), mR
 XM_379625 Homo sapiens hypothetical gene supported by BC048267; NM_178448 (LOC
 XM_379627 Homo sapiens LOC401496 (LOC401496), mRNA
 XM_379628 Homo sapiens LOC401499 (LOC401499), mRNA
 XM_379629 Homo sapiens hypothetical gene supported by AL512690 (LOC401500), mR
 XM_379630 Homo sapiens LOC401503 (LOC401503), mRNA
 XM_379632 Homo sapiens hypothetical protein LOC158376 (LOC158376), mRNA
 XM_379634 Homo sapiens hypothetical gene supported by AK091718 (LOC401504), mR
 XM_379635 Homo sapiens LOC401506 (LOC401506), mRNA
 XM_379636 Homo sapiens hypothetical protein LOC158228 (LOC158228), mRNA
 XM_379637 Homo sapiens hypothetical gene supported by AK127732 (LOC401513), mR
 XM_379638 Homo sapiens hypothetical gene supported by AK026419 (LOC401518), mR
 XM_379639 Homo sapiens LOC401522 (LOC401522), mRNA
 XM_379640 Homo sapiens hypothetical gene supported by BC044751; NM_175923 (LOC
 XM_379641 Homo sapiens hypothetical gene supported by BC062724 (LOC401527), mR
 XM_379642 Homo sapiens hypothetical gene supported by BC032955 (LOC401528), mR
 XM_379643 Homo sapiens hypothetical gene supported by BC032955 (LOC401530), mR
 XM_379644 Homo sapiens hypothetical gene supported by BC000228 (LOC401532), mR
 XM_379645 Homo sapiens hypothetical gene supported by AK094988 (LOC401536), mR
 XM_379647 Homo sapiens similar to hypothetical protein LOC286238 (LOC401538), mR
 XM_379648 Homo sapiens hypothetical gene supported by AK124333 (LOC401539), mR
 XM_379650 Homo sapiens hypothetical protein LOC340515 (LOC340515), mRNA
 XM_379651 Homo sapiens chromosome 9 open reading frame 44 (C9orf44), mRNA
 XM_379655 Homo sapiens hypothetical gene supported by BC031959 (LOC401542), mR
 XM_379656 Homo sapiens hypothetical gene supported by AK092137 (LOC401543), mR
 XM_379657 Homo sapiens hypothetical gene supported by BC043559 (LOC401544), mR
 XM_379660 Homo sapiens LOC401545 (LOC401545), mRNA
 XM_379664 Homo sapiens hypothetical gene supported by BX647840 (LOC401549), mR
 XM_379665 Homo sapiens hypothetical protein LOC286333 (LOC286333), mRNA
 XM_379667 Homo sapiens hypothetical gene supported by BC039180 (LOC401550), mR
 XM_379668 Homo sapiens hypothetical gene supported by AK124723; AL833509 (LOC4
 XM_379671 Homo sapiens hypothetical gene supported by BC019073; BC036842; BC04
 XM_379672 Homo sapiens hypothetical gene supported by AK128673 (LOC401554), mR
 XM_379676 Homo sapiens hypothetical gene supported by AK127261 (LOC401557), mR
 XM_379677 Homo sapiens hypothetical gene supported by AK094119 (LOC401558), mR
 XM_379678 Homo sapiens hypothetical gene supported by BC029166 (LOC401559), mR
 XM_379680 Homo sapiens hypothetical gene supported by AK023162 (LOC90120), mR
 XM_379682 Homo sapiens hypothetical gene supported by AY129027 (LOC401561), mR
 XM_379684 Homo sapiens hypothetical protein LOC286238 (LOC286238), mRNA
 XM_379686 Homo sapiens hypothetical gene supported by AK074437 (LOC401571), mR
 XM_379688 Homo sapiens hypothetical gene supported by BX648912 (LOC401573), mR
 XM_379690 Homo sapiens hypothetical protein LOC284593 (LOC284593), mRNA

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XM_379691 Homo sapiens hypothetical protein LOC284600 (LOC284600), mRNA
 XM_379692 Homo sapiens LOC401574 (LOC401574), mRNA
 XM_379693 Homo sapiens LOC401575 (LOC401575), mRNA
 XM_379694 Homo sapiens hypothetical gene supported by AK125149 (LOC401577), mR
 XM_379695 Homo sapiens hypothetical gene supported by BC056508; NM_004679 (LOC
 XM_379696 Homo sapiens hypothetical gene supported by AK057918 (LOC401579), mR
 XM_379697 Homo sapiens hypothetical gene supported by AL832542 (LOC401582), mR
 XM_379699 Homo sapiens LOC401583 (LOC401583), mRNA
 XM_379700 Homo sapiens hypothetical protein LOC286442 (LOC286442), mRNA
 XM_379702 Homo sapiens hypothetical gene supported by AK098783 (LOC401585), mR
 XM_379703 Homo sapiens hypothetical gene supported by AK130892 (LOC401587), mR
 XM_379704 Homo sapiens hypothetical gene supported by AK056314; BC034616 (LOC4
 XM_379705 Homo sapiens hypothetical protein LOC158572 (LOC158572), mRNA
 XM_379714 Homo sapiens LOC401596 (LOC401596), mRNA
 XM_379715 Homo sapiens hypothetical gene supported by AK125301 (LOC401597), mR
 XM_379716 Homo sapiens hypothetical gene supported by AK057748 (LOC401599), mR
 XM_379717 Homo sapiens hypothetical gene supported by AJ421269; AL359612; NM_0
 XM_379720 Homo sapiens hypothetical gene supported by BX537697 (LOC401613), mR
 XM_379721 Homo sapiens hypothetical gene supported by BX640956 (LOC401615), mR
 XM_379722 Homo sapiens hypothetical gene supported by AK057519 (LOC401616), mR
 XM_379723 Homo sapiens hypothetical gene supported by AK094280 (LOC401617), mR
 XM_379728 Homo sapiens hypothetical protein LOC286411 (LOC286411), mRNA
 XM_379730 Homo sapiens hypothetical gene supported by BC010531 (LOC401621), mR
 XM_379735 Homo sapiens LOC401628 (LOC401628), mRNA
 XM_379736 Homo sapiens hypothetical gene supported by AK125149 (LOC401628), mR
 XM_379738 Homo sapiens LOC401629 (LOC401629), mRNA
 XM_379739 Homo sapiens LOC401630 (LOC401630), mRNA
 XM_379741 Homo sapiens LOC401701 (LOC401701), mRNA
 XM_379749 Homo sapiens LOC401880 (LOC401880), mRNA
 XM_379760 Homo sapiens LOC402387 (LOC402387), mRNA
 XM_379761 Homo sapiens LOC402433 (LOC402433), mRNA
 XM_379766 Homo sapiens unc-84 homolog A (C. elegans) (UNC84A), mRNA
 XM_379767 Homo sapiens DKFZP586J0619 protein (DKFZP586J0619), mRNA
 XM_379771 Homo sapiens KIAA0415 gene product (KIAA0415), mRNA
 XM_379772 Homo sapiens hypothetical gene supported by BC031661 (LOC402450), mR
 XM_379773 Homo sapiens hypothetical protein LOC285924 (LOC285924), mRNA
 XM_379774 Homo sapiens KIAA1856 protein (KIAA1856), mRNA
 XM_379775 Homo sapiens hypothetical gene supported by AK125308 (LOC402452), mR
 XM_379776 Homo sapiens hypothetical gene supported by AK123535 (LOC402454), mR
 XM_379777 Homo sapiens similar to Oncomodulin (OM) (Parvalbumin beta) (LOC402456
 XM_379780 Homo sapiens similar to unc-93 homolog B1; unc93 (C.elegans) homolog B;
 XM_379781 Homo sapiens similar to beta-1,4-mannosyltransferase; beta-1,4 mannosyltr
 XM_379783 Homo sapiens hypothetical gene supported by AK027125 (LOC402460), mR
 XM_379784 Homo sapiens glucocorticoid induced transcript 1 (GLCCH1), mRNA
 XM_379786 Homo sapiens similar to Chain , Heat-Shock Cognate 70kd Protein (44kd At
 XM_379787 Homo sapiens similar to heat shock 70kDa protein 8 Isoform 2; heat shock c
 XM_379788 Homo sapiens PHD finger protein 14 (PHF14), mRNA
 XM_379792 Homo sapiens similar to TWIST neighbor (LOC402464), mRNA
 XM_379793 Homo sapiens ribosomal protein L21 (RPL21), mRNA
 XM_379794 Homo sapiens similar to ribosomal protein L23 (LOC402465), mRNA
 XM_379796 Homo sapiens similar to Dual specificity protein kinase CLK2 (CDC like kinas
 XM_379797 Homo sapiens similar to FKSG54 (LOC402469), mRNA
 XM_379798 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA
 XM_379800 Homo sapiens KIAA0644 gene product (KIAA0644), mRNA
 XM_379801 Homo sapiens KIAA0241 protein (KIAA0241), mRNA
 XM_379802 Homo sapiens LOC89231 (LOC89231), mRNA
 XM_379803 Homo sapiens similar to KIAA0877 protein (LOC402477), mRNA

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XM_379806 Homo sapiens similar to RP9 protein (LOC402478), mRNA
 XM_379807 Homo sapiens similar to RIKEN cDNA 9330128H10 gene (LOC402479), mR
 XM_379809 Homo sapiens hypothetical gene supported by AF447883 (LOC402481), mR
 XM_379812 Homo sapiens similar to TRGV9 (LOC402482), mRNA
 XM_379815 Homo sapiens similar to sequence-specific single-stranded-DNA-binding pro
 XM_379816 Homo sapiens similar to t-complex 1; T-complex locus TCP-1; t-complex 1 (a
 XM_379817 Homo sapiens similar to RAS p21 protein activator 4; GTPase activating prot
 XM_379818 Homo sapiens similar to RAS p21 protein activator 4; GTPase activating prot
 XM_379819 Homo sapiens similar to DNA directed RNA polymerase II polypeptide J-retai
 XM_379820 Homo sapiens similar to Ras GTPase-activating protein 4 (RasGAP-activatin
 XM_379824 Homo sapiens similar to cell division cycle 10 homolog (LOC402491), mRNA
 XM_379825 Homo sapiens similar to hypothetical protein FLJ25976 (LOC402492), mRN/
 XM_379827 Homo sapiens hypothetical gene supported by AK126096 (LOC402494), mR
 XM_379830 Homo sapiens similar to KIAA0207 (LOC402495), mRNA
 XM_379831 Homo sapiens hypothetical gene supported by AK097404; NM_198284 (LOC
 XM_379832 Homo sapiens hypothetical gene supported by AK127870 (LOC402497), mR
 XM_379834 Homo sapiens similar to SMT3 suppressor of mif two 3 homolog 2 (LOC4025
 XM_379835 Homo sapiens similar to hypothetical protein DKFZp434F142 (LOC402501),
 XM_379836 Homo sapiens similar to septin 10 isoform 1 (LOC402502), mRNA
 XM_379838 Homo sapiens similar to bA145E8.1 (KIAA1074) (LOC402505), mRNA
 XM_379839 Homo sapiens similar to CAGL79 (LOC402508), mRNA
 XM_379840 Homo sapiens similar to solute carrier family 29 (nucleoside transporters), m
 XM_379841 Homo sapiens zinc finger protein 479 (ZNF479), mRNA
 XM_379842 Homo sapiens LOC402513 (LOC402513), mRNA
 XM_379843 Homo sapiens hypothetical gene supported by BC040831 (LOC402514), mR
 XM_379844 Homo sapiens hypothetical gene supported by BC040831 (LOC402517), mR
 XM_379845 Homo sapiens similar to solute carrier family 29 (nucleoside transporters), m
 XM_379846 Homo sapiens similar to BC060615 protein (LOC402519), mRNA
 XM_379847 Homo sapiens similar to CAGL79 (LOC402520), mRNA
 XM_379848 Homo sapiens similar to hypothetical protein LOC285908 (LOC402521), mR/
 XM_379849 Homo sapiens similar to GA binding protein transcription factor, alpha subun
 XM_379850 Homo sapiens similar to hypothetical ZNF-like protein (LOC402524), mRNA
 XM_379851 Homo sapiens similar to RPL6 protein (LOC402525), mRNA
 XM_379852 Homo sapiens similar to envelope protein (LOC402526), mRNA
 XM_379853 Homo sapiens similar to myelin protein zero-like 1; protein zero related (LOC
 XM_379854 Homo sapiens similar to hypothetical protein FLJ25037 (LOC402529), mR
 XM_379855 Homo sapiens similar to hypothetical protein MGC16733 similar to CG12113
 XM_379856 Homo sapiens similar to MGC16733 protein (LOC402531), mRNA
 XM_379857 Homo sapiens similar to 60S ribosomal protein L35 (LOC402536), mRNA
 XM_379858 Homo sapiens similar to Williams-Beuren syndrome critical region protein 19
 XM_379859 Homo sapiens similar to hypothetical protein FLJ10900 (LOC402540), mRN/
 XM_379860 Homo sapiens similar to Williams Beuren syndrome chromosome region 19 i
 XM_379861 Homo sapiens similar to FKBP6 protein (LOC402543), mRNA
 XM_379862 Homo sapiens similar to Brutons tyrosine kinase-associated protein-135; BA/
 XM_379863 Homo sapiens similar to Neutrophil cytosol factor 1 (NCF-1) (Neutrophil NAD
 XM_379864 Homo sapiens similar to transcription factor GTF2IRD2 (LOC402546), mRNA
 XM_379865 Homo sapiens similar to Nuclear envelope pore membrane protein POM 121
 XM_379866 Homo sapiens Williams Beuren syndrome chromosome region 24 (WBSCR2
 XM_379868 Homo sapiens similar to Neutrophil cytosolic factor 1 (LOC402549), mRNA
 XM_379869 Homo sapiens similar to transcription factor GTF2IRD2 (LOC402550), mRNA/
 XM_379871 Homo sapiens similar to RCC1-like G exchanging factor-like isoform 1; RCC-
 XM_379872 Homo sapiens similar to PMS4 (LOC402552), mRNA
 XM_379874 Homo sapiens similar to PMS4 homolog mismatch repair protein - human (Lt
 XM_379875 Homo sapiens similar to FKBP6 protein (LOC402555), mRNA
 XM_379876 Homo sapiens tripartite motif-containing 50B (TRIM50B), mRNA
 XM_379877 Homo sapiens similar to Nuclear envelope pore membrane protein POM 121
 XM_379879 Homo sapiens similar to hypothetical protein LOC285908 (LOC402558), mR/

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XM_379881 Homo sapiens similar to hypothetical protein (LOC402559), mRNA
 XM_379883 Homo sapiens similar to Piccolo protein (Aczonin) (LOC402561), mRNA
 XM_379884 Homo sapiens sema domain, immunoglobulin domain (Ig), short basic doma
 XM_379885 Homo sapiens similar to Heterogeneous nuclear ribonucleoprotein A1 (Helix-
 XM_379887 Homo sapiens similar to hypothetical protein 4932412H11 (LOC402565), mF
 XM_379891 Homo sapiens hypothetical gene supported by AK124274 (LOC402568), mR
 XM_379892 Homo sapiens similar to Homeobox protein DLX-6 (LOC402567), mRNA
 XM_379893 Homo sapiens similar to CG14980-PB (LOC375601), mRNA
 XM_379894 Homo sapiens similar to importin alpha 1b (LOC402569), mRNA
 XM_379895 Homo sapiens hypothetical protein FLJ22037 (FLJ22037), mRNA
 XM_379896 Homo sapiens hypothetical protein LOC285989 (LOC285989), mRNA
 XM_379897 Homo sapiens similar to Zinc-alpha-2-glycoprotein precursor (Zn-alpha-2-gly
 XM_379899 Homo sapiens similar to hypothetical protein MGC49416 (LOC402572), mRN
 XM_379901 Homo sapiens hypothetical gene supported by BC031966 (LOC402573), mR
 XM_379904 Homo sapiens similar to mucin 11 (LOC402575), mRNA
 XM_379905 Homo sapiens similar to intestinal membrane mucin MUC17 (LOC402576), n
 XM_379906 Homo sapiens similar to PMS5 homolog mismatch repair protein - human (L
 XM_379908 Homo sapiens similar to DNA directed RNA polymerase II polypeptide J-rela
 XM_379909 Homo sapiens similar to M-phase phosphoprotein 11 (LOC402580), mRNA
 XM_379910 Homo sapiens similar to reverse transcriptase related protein (LOC402581),
 XM_379911 Homo sapiens similar to KIAA1218 protein (LOC402584), mRNA
 XM_379913 Homo sapiens similar to calcium-independent phospholipase A2 (LOC40258
 XM_379914 Homo sapiens hypothetical protein LOC288009 (LOC288009), mRNA
 XM_379917 Homo sapiens similar to hyaluronoglucosaminidase 1 Isoform 1; hyaluronida
 XM_379919 Homo sapiens similar to hypothetical protein FLJ25976 (LOC402594), mRN/
 XM_379920 Homo sapiens similar to hypothetical protein LOC285908 (LOC402595), mR/
 XM_379921 Homo sapiens hypothetical protein LOC346653 (LOC346653), mRNA
 XM_379923 Homo sapiens KIAA1170 protein (KIAA1170), mRNA
 XM_379927 Homo sapiens plexin A4 (PLXNA4), mRNA
 XM_379931 Homo sapiens hypothetical protein LOC155006 (LOC155006), mRNA
 XM_379932 Homo sapiens KIAA1549 protein (KIAA1549), mRNA
 XM_379933 Homo sapiens hypothetical protein FLJ25778 (FLJ25778), mRNA
 XM_379934 Homo sapiens similar to A630082K20Rik protein (LOC402600), mRNA
 XM_379935 Homo sapiens similar to RAB19, member RAS oncogene family (LOC40260
 XM_379936 Homo sapiens similar to hypothetical protein (LOC402602), mRNA
 XM_379938 Homo sapiens LCHN protein (LCHN), mRNA
 XM_379939 Homo sapiens similar to RIKEN cDNA 1700018G05 (LOC402604), mRNA
 XM_379940 Homo sapiens hypothetical protein LOC93432 (LOC93432), mRNA
 XM_379954 Homo sapiens similar to KIAA0738 protein (LOC402618), mRNA
 XM_379956 Homo sapiens similar to hypothetical protein MGC41943 (LOC402619), mR
 XM_379959 Homo sapiens FLJ43692 protein (FLJ43692), mRNA
 XM_379962 Homo sapiens similar to KIAA1285 protein (LOC402620), mRNA
 XM_379964 Homo sapiens likely ortholog of mouse zinc finger protein EZI (EZI), mRNA
 XM_379965 Homo sapiens FLJ45737 protein (FLJ45737), mRNA
 XM_379966 Homo sapiens similar to KIAA2036 protein (LOC402621), mRNA
 XM_379967 Homo sapiens KIAA0543 protein (KIAA0543), mRNA
 XM_379968 Homo sapiens chromosome 7 open reading frame 32 (C7orf32), mRNA
 XM_379970 Homo sapiens similar to Zinc finger protein 84 (Zinc finger protein HPF2) (LC
 XM_379974 Homo sapiens KIAA1402 protein (CSG1A-T), mRNA
 XM_379975 Homo sapiens hypothetical gene supported by AK127717 (LOC402625), mR
 XM_379976 Homo sapiens similar to Fatty acid-binding protein, epidermal (E-FABP) (Psc
 XM_379977 Homo sapiens similar to T-complex protein 1 (LOC402629), mRNA
 XM_379979 Homo sapiens hypothetical protein LOC285888 (LOC285888), mRNA
 XM_379980 Homo sapiens hypothetical protein LOC155435 (LOC155435), mRNA
 XM_379983 Homo sapiens ubiquitin-protein isopeptide ligase (E3) (KIAA0010), mRNA
 XM_379986 Homo sapiens similar to hypothetical protein FLJ37300 (LOC402633), mRN/
 XM_379987 Homo sapiens similar to mesenchymal stem cell protein DSC92; neurite outg

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XM_379988 Homo sapiens similar to galectin-related inter-fiber protein (LOC402635), mRNA
 XM_379989 Homo sapiens similar to olfactory receptor MOR267-3 (LOC402636), mRNA
 XM_379990 Homo sapiens similar to Calgizzarin (S100C protein) (MLN 70) (LOC402637)
 XM_379991 Homo sapiens similar to ribosomal protein L31 (LOC402638), mRNA
 XM_379993 Homo sapiens similar to olfactory receptor MOR145-2 (LOC402639), mRNA
 XM_379994 Homo sapiens similar to 4930579E17Rik protein (LOC402640), mRNA
 XM_379995 Homo sapiens similar to MGC76216 protein (LOC402641), mRNA
 XM_379996 Homo sapiens similar to solute carrier family 40 (iron-regulated transporter),
 XM_379997 Homo sapiens similar to tropomyosin 3 (LOC402643), mRNA
 XM_379998 Homo sapiens similar to peptidylprolyl isomerase A (LOC402644), mRNA
 XM_379999 Homo sapiens similar to FLJ40113 protein (LOC402645), mRNA
 XM_380002 Homo sapiens similar to ELK1 (LOC402648), mRNA
 XM_380005 Homo sapiens similar to equilibrative nucleoside transporter 4; hENT4 (LOC402662), mRNA
 XM_380006 Homo sapiens similar to beta-glucuronidase (LOC402662), mRNA
 XM_380007 Homo sapiens similar to vomeronasal receptor V1RC-3 (LOC402663), mRNA
 XM_380008 Homo sapiens similar to Opioid binding protein/cell adhesion molecule precu
 XM_380009 Homo sapiens similar to protein kinase related to Raf protein kinases; Methio
 XM_380010 Homo sapiens similar to metabotropic glutamate receptor 8; G protein-couple
 XM_380011 Homo sapiens similar to GTF2I repeat domain containing 1 isoform 2; Williar
 XM_380012 Homo sapiens similar to opposite strand transcription unit to Stag3; Gats pro
 XM_380013 Homo sapiens similar to GrpE protein homolog 1, mitochondrial precursor (V
 XM_380014 Homo sapiens similar to GTF2I repeat domain containing 1 isoform 2; Williar
 XM_380015 Homo sapiens similar to peptidylprolyl isomerase A (LOC402673), mRNA
 XM_380018 Homo sapiens hypothetical protein DKFZP434A0225 (DKFZP434A0225), m
 XM_380019 Homo sapiens similar to RIKEN cDNA 4930511M11 (LOC402675), mRNA
 XM_380020 Homo sapiens similar to Protein C8orf66 (HSPC125) (Myo13 protein) (LOC402675), mRNA
 XM_380021 Homo sapiens similar to ribosomal protein S3a; 40S ribosomal protein S3a; \n
 XM_380022 Homo sapiens similar to Ser/Thr protein kinase PAR-1Balpha (LOC402679),
 XM_380024 Homo sapiens similar to opposite strand transcription unit to Stag3; Gats pro
 XM_380025 Homo sapiens similar to PMS2L13 (LOC402681), mRNA
 XM_380026 Homo sapiens similar to RIKEN cDNA 2700038N03 (LOC402682), mRNA
 XM_380028 Homo sapiens similar to 40S RIBOSOMAL PROTEIN SA (P40) (34/67 KD L/
 XM_380031 Homo sapiens similar to KIAA0538 protein (LOC402684), mRNA
 XM_380032 Homo sapiens similar to Ras GTPase-activating protein 4 (RasGAP-activatin
 XM_380033 Homo sapiens similar to 60S ribosomal protein L23a (LOC402686), mRNA
 XM_380034 Homo sapiens similar to Argininosuccinate synthase (Citrulline--aspartate lig
 XM_380036 Homo sapiens similar to RIKEN cDNA 6332401O19 gene (LOC402689), mR
 XM_380038 Homo sapiens similar to Triosephosphate isomerase (TIM) (LOC402691), m
 XM_380039 Homo sapiens similar to aldo-keto reductase family 1 , member B10; aldose r
 XM_380040 Homo sapiens similar to beta-tubulin (LOC402693), mRNA
 XM_380042 Homo sapiens similar to 60S ribosomal protein L15 (LOC402694), mRNA
 XM_380044 Homo sapiens similar to 60S ribosomal protein L17 (L23) (Amino acid starva
 XM_380045 Homo sapiens similar to Nucleoside diphosphate kinase, mitochondrial precu
 XM_380047 Homo sapiens similar to Nucleoside diphosphate kinase, mitochondrial precu
 XM_380048 Homo sapiens similar to Olfactory receptor 9A4 (LOC402698), mRNA
 XM_380055 Homo sapiens similar to Olfactory receptor 9A2 (LOC402707), mRNA
 XM_380056 Homo sapiens similar to Olfactory receptor 6V1 (LOC402708), mRNA
 XM_380057 Homo sapiens similar to Histidine triad nucleotide-binding protein 1 (Adenosi
 XM_380059 Homo sapiens similar to Olfactory receptor 2F2 (Olfactory receptor 7-1) (OR)
 XM_380063 Homo sapiens similar to Olfactory receptor 2A12 (LOC402711), mRNA
 XM_380067 Homo sapiens similar to Olfactory receptor 2A1 (LOC402712), mRNA
 XM_380068 Homo sapiens similar to Olfactory receptor 2A1 (LOC402713), mRNA
 XM_380069 Homo sapiens similar to OG-2 homeodomain protein-like; similar to U65067
 XM_380070 Homo sapiens similar to Importin alpha-2 subunit (Karyopherin alpha-2 subu
 XM_380072 Homo sapiens similar to 60S ribosomal protein L32 (LOC402716), mRNA
 XM_380073 Homo sapiens similar to Huntingtin interacting protein K (LOC402717), mRNA
 XM_380074 Homo sapiens similar to BET1 homolog (Golgi vesicular membrane traffickin

XM_380076 Homo sapiens similar to TSH receptor suppressor element-binding protein-1
 XM_380077 Homo sapiens similar to ppg3 (LOC402720), mRNA
 XM_380078 Homo sapiens similar to S-adenosylmethionine decarboxylase 1; S-adenosyl
 XM_380081 Homo sapiens hypothetical protein FLJ36112 (FLJ36112), mRNA
 XM_380082 Homo sapiens hypothetical gene supported by AK125766 (LOC402448), mR
 XM_380085 Homo sapiens hypothetical gene supported by BC032734 (LOC402449), mR
 XM_380087 Homo sapiens LOC402451 (LOC402451), mRNA
 XM_380088 Homo sapiens LOC402453 (LOC402453), mRNA
 XM_380089 Homo sapiens hypothetical gene supported by AK127500 (LOC402463), mR
 XM_380091 Homo sapiens hypothetical gene supported by BC039682 (LOC402466), mR
 XM_380092 Homo sapiens hypothetical protein LOC285941 (LOC285941), mRNA
 XM_380093 Homo sapiens hypothetical gene supported by BC025338 (LOC402470), mR
 XM_380094 Homo sapiens hypothetical gene supported by AK093987 (LOC402471), mR
 XM_380095 Homo sapiens LOC402472 (LOC402472), mRNA
 XM_380096 Homo sapiens LOC402473 (LOC402473), mRNA
 XM_380097 Homo sapiens hypothetical gene supported by BC023581; BC044638 (LOC4
 XM_380098 Homo sapiens hypothetical gene supported by BC016976 (LOC402475), mR
 XM_380099 Homo sapiens hypothetical gene supported by AK092714 (LOC402476), mR
 XM_380100 Homo sapiens hypothetical gene supported by AL832082 (LOC402480), mR
 XM_380103 Homo sapiens hypothetical gene supported by AK127273 (LOC402483), mR
 XM_380104 Homo sapiens LOC402485 (LOC402485), mRNA
 XM_380105 Homo sapiens hypothetical protein LOC285958 (LOC285958), mRNA
 XM_380106 Homo sapiens hypothetical gene supported by AK125311 (LOC402493), mR
 XM_380107 Homo sapiens hypothetical gene supported by AK024248; AL137733; BC06;
 XM_380108 Homo sapiens LOC402506 (LOC402506), mRNA
 XM_380109 Homo sapiens LOC402507 (LOC402507), mRNA
 XM_380110 Homo sapiens LOC402511 (LOC402511), mRNA
 XM_380111 Homo sapiens hypothetical gene supported by BX646489 (LOC402512), mR
 XM_380112 Homo sapiens LOC402515 (LOC402515), mRNA
 XM_380113 Homo sapiens LOC402516 (LOC402516), mRNA
 XM_380114 Homo sapiens LOC402523 (LOC402523), mRNA
 XM_380115 Homo sapiens LOC402528 (LOC402528), mRNA
 XM_380117 Homo sapiens LOC402532 (LOC402532), mRNA
 XM_380118 Homo sapiens hypothetical gene supported by AK024371; BC037920 (LOC4
 XM_380119 Homo sapiens hypothetical gene supported by AK054923; AK126730; NM_0
 XM_380120 Homo sapiens LOC402537 (LOC402537), mRNA
 XM_380121 Homo sapiens LOC402538 (LOC402538), mRNA
 XM_380122 Homo sapiens LOC402542 (LOC402542), mRNA
 XM_380125 Homo sapiens LOC402553 (LOC402553), mRNA
 XM_380126 Homo sapiens hypothetical gene supported by AK091784 (LOC402557), mR
 XM_380127 Homo sapiens LOC402560 (LOC402560), mRNA
 XM_380128 Homo sapiens LOC402563 (LOC402563), mRNA
 XM_380129 Homo sapiens LOC402564 (LOC402564), mRNA
 XM_380131 Homo sapiens hypothetical protein FLJ34048 (FLJ34048), mRNA
 XM_380133 Homo sapiens LOC402568 (LOC402568), mRNA
 XM_380134 Homo sapiens LOC402574 (LOC402574), mRNA
 XM_380135 Homo sapiens LOC402578 (LOC402578), mRNA
 XM_380136 Homo sapiens LOC402582 (LOC402582), mRNA
 XM_380137 Homo sapiens hypothetical protein LOC340340 (LOC340340), mRNA
 XM_380138 Homo sapiens LOC402586 (LOC402586), mRNA
 XM_380139 Homo sapiens hypothetical gene supported by BX537645 (LOC402587), mR
 XM_380140 Homo sapiens LOC402589 (LOC402589), mRNA
 XM_380141 Homo sapiens hypothetical gene supported by BX648695 (LOC402590), mR
 XM_380142 Homo sapiens hypothetical gene supported by BC063892 (LOC402591), mR
 XM_380143 Homo sapiens LOC402592 (LOC402592), mRNA
 XM_380146 Homo sapiens hypothetical gene supported by AK125651; BC039420 (LOC4
 XM_380147 Homo sapiens LOC402597 (LOC402597), mRNA

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XM_380148 Homo sapiens LOC402598 (LOC402598), mRNA
XM_380149 Homo sapiens LOC402599 (LOC402599), mRNA
XM_380150 Homo sapiens hypothetical gene supported by BX648692 (LOC402603), mR
XM_380151 Homo sapiens LOC402617 (LOC402617), mRNA
XM_380152 Homo sapiens hypothetical protein LOC90520 (LOC90520), mRNA
XM_380153 Homo sapiens LOC402622 (LOC402622), mRNA
XM_380154 Homo sapiens hypothetical protein LOC285972 (LOC285972), mRNA
XM_380155 Homo sapiens LOC402626 (LOC402626), mRNA
XM_380156 Homo sapiens LOC402627 (LOC402627), mRNA
XM_380157 Homo sapiens LOC402630 (LOC402630), mRNA
XM_380158 Homo sapiens hypothetical gene supported by AY166699 (LOC402631), mR
XM_380159 Homo sapiens hypothetical gene supported by AK054822 (LOC402632), mR
XM_380160 Homo sapiens hypothetical protein LOC285889 (LOC285889), mRNA
XM_380162 Homo sapiens hypothetical protein LOC154822 (LOC154822), mRNA
XM_380170 Homo sapiens phosphodiesterase 4D interacting protein (myomegalin) (PDE
XM_380171 Homo sapiens iroquois homeobox protein 1 (IRX1), mRNA
XM_380173 Homo sapiens hypothetical protein MGC39830 (MGC39830), mRNA
XM_380174 Homo sapiens similar to aquaporin 11 (LOC285192), mRNA
XM_380175 Homo sapiens hypothetical protein MGC22265 (MGC22265), mRNA
XM_380176 Homo sapiens similar to hypothetical protein D030010E02 (LOC401952), mF
NC_000001 Homo sapiens chromosome 1, complete sequence
NC_000002 Homo sapiens chromosome 2, complete sequence
NC_000003 Homo sapiens chromosome 3, complete sequence
NC_000004 Homo sapiens chromosome 4, complete sequence
NC_000005 Homo sapiens chromosome 5, complete sequence
NC_000006 Homo sapiens chromosome 6, complete sequence
NC_000007 Homo sapiens chromosome 7, complete sequence
NC_000008 Homo sapiens chromosome 8, complete sequence
NC_000009 Homo sapiens chromosome 9, complete sequence
NC_000010 Homo sapiens chromosome 10, complete sequence
NC_000011 Homo sapiens chromosome 11, complete sequence
NC_000012 Homo sapiens chromosome 12, complete sequence
NC_000013 Homo sapiens chromosome 13, complete sequence
NC_000014 Homo sapiens chromosome 14, complete sequence
NC_000015 Homo sapiens chromosome 15, complete sequence
NC_000016 Homo sapiens chromosome 16, complete sequence
NC_000017 Homo sapiens chromosome 17, complete sequence
NC_000018 Homo sapiens chromosome 18, complete sequence
NC_000019 Homo sapiens chromosome 19, complete sequence
NC_000020 Homo sapiens chromosome 20, complete sequence
NC_000021 Homo sapiens chromosome 21, complete sequence
NC_000022 Homo sapiens chromosome 22, complete sequence
NC_000023 Homo sapiens chromosome X, complete sequence
NC_000024 Homo sapiens chromosome Y, complete sequence
NC_001807 Homo sapiens mitochondrion, complete genome
NG_000002 Homo sapiens immunoglobulin lambda locus (IGL@) on chromosome 22
NG_000004 Homo sapiens cytochrome P450, family 3, subfamily A (CYP3A) on chromos
NG_000006 Homo sapiens alpha globin region (HBA@) on chromosome 16
NG_000007 Homo sapiens beta globin region (HBB@) on chromosome 11
NG_000008 Homo sapiens cytochrome P450, family 2, subfamily A (CYP2A) on chromos
NG_000009 Homo sapiens small histone family cluster (HFS@) on chromosome 6
NG_000012 Homo sapiens protocadherin gamma cluster (PCDHG@) on chromosome 5
NG_000013 Homo sapiens MHC class III complement gene cluster, monomodular haplot
NG_000016 Homo sapiens genomic protocadherin alpha cluster (PCDHA@) on chromos
NG_000017 Homo sapiens protocadherin beta cluster (PCDHB@) on chromosome 5
NG_000018 Homo sapiens type I (acidic) hair keratin gene cluster (KRTHA.1@) on chro
NG_000019 Homo sapiens chorionic gonadotropin beta region (CGB@) on chromosome

NG_000827 Homo sapiens genomic histone family microcluster (HFM@) on chromosome
 NG_000833 Homo sapiens immunoglobulin kappa locus, distal duplicated V-cluster (IGK-
 NG_000834 Homo sapiens immunoglobulin kappa locus, proximal V-cluster and J-C clusl
 NG_000837 Homo sapiens surfet locus (SURF@) on chromosome 9
 NG_000839 Homo sapiens cystatin locus (CST@) on chromosome 20
 NG_000840 Homo sapiens actin, beta pseudogene 8 (ACTBP8) on chromosome 6
 NG_000841 Homo sapiens actin, gamma pseudogene 2 (ACTGP2) on chromosome Y
 NG_000842 Homo sapiens actin, gamma pseudogene 3 (ACTGP3) on chromosome 20
 NG_000843 Homo sapiens adenosine A2b receptor pseudogene (ADORA2BP) on chrom
 NG_000845 Homo sapiens argininosuccinate synthetase pseudogene 2 (ASSP2) on chro
 NG_000846 Homo sapiens argininosuccinate synthetase pseudogene 4 (ASSP4) on chro
 NG_000847 Homo sapiens argininosuccinate synthetase pseudogene 5 (ASSP5) on chro
 NG_000848 Homo sapiens argininosuccinate synthetase pseudogene 6 (ASSP6) on chro
 NG_000849 Homo sapiens ATPase, Na+/K+ transporting, beta 3 pseudogene (ATP1B3P
 NG_000850 Homo sapiens BCL2-like 7 pseudogene 1 (BCL2L7P1) on chromosome 20
 NG_000851 Homo sapiens crystallin, beta B2 pseudogene 1 (CRYBB2P1) on chromoson
 NG_000852 Homo sapiens cysteine and glycine-rich protein 2 pseudogene (CSR2P2) on
 NG_000853 Homo sapiens cytochrome P450, family 2, subfamily D, polypeptide 8 pseud
 NG_000854 Homo sapiens cytochrome P450, family 2, subfamily D, polypeptide 8 pseud
 NG_000858 Homo sapiens dihydrofolate reductase pseudogene 1 (DHFRP1) on chromo
 NG_000859 Homo sapiens eukaryotic translation initiation factor 5A pseudogene 1 (EIF5
 NG_000861 Homo sapiens glycerol kinase pseudogene 3 (GKP3) on chromosome 4
 NG_000862 Homo sapiens quinoline nucleotide binding protein (G protein), q polypeptide |
 NG_000863 Homo sapiens hydroxyacyl-Coenzyme A dehydrogenase-3/ketoacyl-Coenzym
 NG_000865 Homo sapiens proliferating cell nuclear antigen pseudogene (PCNAP) on chr
 NG_000866 Homo sapiens RNA binding motif, single stranded interacting protein 1, pseu
 NG_000867 Homo sapiens radixin pseudogene 2 (RDXP2) on chromosome X
 NG_000868 Homo sapiens ribosomal protein L21 pseudogene 1 (RPL21P1) on chromos
 NG_000869 Homo sapiens ribosomal protein L32 pseudogene 1 (RPL32P1) on chromos
 NG_000870 Homo sapiens ribonucleotide reductase M2 polypeptide pseudogene 3 (RR1/
 NG_000871 Homo sapiens ribonucleotide reductase M2 polypeptide pseudogene 4 (RR1/
 NG_000872 Homo sapiens tRNA phosphoserine (opal suppressor) pseudogene 1 (TRSP
 NG_000873 Homo sapiens makorin, ring finger protein, pseudogene 1 (MKRNP1) on chr
 NG_000874 Homo sapiens heterogeneous nuclear ribonucleoprotein D (AU-rich element
 NG_000877 Homo sapiens ribosomal protein L3 pseudogene 1 (RPL3P1) on chromosom
 NG_000878 Homo sapiens ribosomal protein L23a pseudogene 3 (RPL23AP3) on chrom
 NG_000880 Homo sapiens arylsulfatase E pseudogene (ARSEP) on chromosome Y
 NG_000881 Homo sapiens arylsulfatase D pseudogene (ARSDP) on chromosome Y
 NG_000882 Homo sapiens voltage-dependent anion channel 5, pseudogene (VDAC5P) c
 NG_000883 Homo sapiens high-mobility group (nonhistone chromosomal) protein 1-like 5
 NG_000884 Homo sapiens snail homolog 1 like 1 (Drosophila) (SNAI1L1) pseudogene o
 NG_000885 Homo sapiens nucleophosmin 1 (nucleolar phosphoprotein B23, numatrin) p
 NG_000886 Homo sapiens nucleophosmin 1 (nucleolar phosphoprotein B23, numatrin) p
 NG_000887 Homo sapiens nucleophosmin 1 (nucleolar phosphoprotein B23, numatrin) p
 NG_000889 Homo sapiens ribosomal protein L41, pseudogene 3 (RPL41P3) on chromos
 NG_000890 Homo sapiens ribosomal protein L41, pseudogene 2 (RPL41P2) on chromos
 NG_000891 Homo sapiens ribosomal protein L41, pseudogene 1 (RPL41P1) on chromos
 NG_000892 Homo sapiens high-mobility group (nonhistone chromosomal) protein 17-like
 NG_000893 Homo sapiens ribosomal protein S24 pseudogene 1 (RPS24P1) on chromos
 NG_000894 Homo sapiens ribosomal protein L34 pseudogene 2 (RPL34P2) on chromos
 NG_000895 Homo sapiens ribosomal protein L34 pseudogene 1 (RPL34P1) on chromos
 NG_000896 Homo sapiens ribosomal protein S5 pseudogene 1 (RPS5P1) on chromoson
 NG_000897 Homo sapiens high-mobility group (nonhistone chromosomal) protein 1-like 5
 NG_000898 Homo sapiens zinc finger protein 299 pseudogene (ZNF299P) on chromosor
 NG_000899 Homo sapiens voltage-dependent anion channel 2 pseudogene (VDAC2P) o
 NG_000900 Homo sapiens tubulin, alpha pseudogene (TUBAP) on chromosome 21
 NG_000901 Homo sapiens solute carrier family 6, member 6 pseudogene (SLC6A6P) on

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NG_000903 Homo sapiens ribosomal protein S5-like (RPS5L) pseudogene on chromosome
 NG_000904 Homo sapiens ribosomal protein S3A pseudogene 1 (RPS3AP1) on chromosome
 NG_000906 Homo sapiens ribosomal protein S20, pseudogene 1 (RPS20P1) on chromosome
 NG_000907 Homo sapiens ribosomal protein L34 pseudogene 3 (RPL34P3) on chromosome
 NG_000908 Homo sapiens ribosomal protein L31 pseudogene 1 (RPL31P1) on chromosome
 NG_000909 Homo sapiens ribosomal protein L23 pseudogene 2 (RPL23P2) on chromosome
 NG_000910 Homo sapiens ribosomal protein L23a pseudogene 4 (RPL23AP4) on chromosome
 NG_000911 Homo sapiens ribosomal protein L10 pseudogene 1 (RPL10P1) on chromosome
 NG_000912 Homo sapiens ribosomal modification protein rimK-like (E. coli) pseudogene
 NG_000913 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 2 pseudogene
 NG_000914 Homo sapiens peptidylprolyl isomerase A (cyclophilin A), pseudogene (PPIA)
 NG_000915 Homo sapiens polymerase (RNA) II (DNA directed) polypeptide C, pseudogene
 NG_000916 Homo sapiens poly(rC) binding protein 2, pseudogene 1 (PCBP2P1) on chromosome
 NG_000917 Homo sapiens myosin, light polypeptide 6, pseudogene (MYL6P) on chromosome
 NG_000919 Homo sapiens inner membrane protein, mitochondrial (mitofilin) pseudogene
 NG_000920 Homo sapiens heat shock 60kDa protein 1 (chaperonin) pseudogene 7 (HSP)
 NG_000921 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 pso
 NG_000922 Homo sapiens H2A histone family, member Z, pseudogene (H2AFZP) on chromosome
 NG_000923 Homo sapiens farnesyl diphosphate synthase pseudogene (FDPSP) on chromosome
 NG_000924 Homo sapiens eukaryotic translation initiation factor 4A, isoform 1, pseudogene
 NG_000925 Homo sapiens eukaryotic translation initiation factor 3, subunit 5 epsilon, 47k
 NG_000927 Homo sapiens cytochrome P450, family 4, subfamily F, polypeptide 3-like ps
 NG_000929 Homo sapiens complement component 1, q subcomponent binding protein, f
 NG_000930 Homo sapiens F-box and WD-40 domain protein 11 pseudogene 1 (FBXW11)
 NG_000933 Homo sapiens endoplasmic reticulum luminal protein 28 pseudogene (ERP)
 NG_000936 Homo sapiens plakophilin 2 pseudogene 1 (PKP2P1) on chromosome 12
 NG_000938 Homo sapiens keratin, hair, basic, pseudogene 1 (KRTHBP1) on chromosome
 NG_000939 Homo sapiens keratin associated protein 2 pseudogene 1 (KRTAP2P1) on chromosome
 NG_000940 Homo sapiens keratin, hair, basic, pseudogene 2 (KRTHBP2) on chromosome
 NG_000941 Homo sapiens keratin associated protein 3 pseudogene 1 (KRTAP3P1) on chromosome
 NG_000942 Homo sapiens keratin associated protein 9 pseudogene 1 (KRTAP9P1) on chromosome
 NG_000943 Homo sapiens keratin, hair, basic, pseudogene 3 (KRTHBP3) on chromosome
 NG_000944 Homo sapiens keratin, hair, basic, pseudogene 4 (KRTHBP4) on chromosome
 NG_000945 Homo sapiens ribosomal protein L12-like 3 (RPL12L3) pseudogene on chromosome
 NG_000946 Homo sapiens ribosomal protein S2-like 1 (RPS2L1) pseudogene on chromosome
 NG_000948 Homo sapiens 5'-nucleotidase, cytosolic III pseudogene 1 (NT5C3P1) on chromosome
 NG_000949 Homo sapiens pleckstrin homology domain containing, family A member 3 ps
 NG_000950 Homo sapiens ribosomal protein S4-like 2 (RPS4L2) pseudogene on chromosome
 NG_000951 Homo sapiens ribosomal protein S15a pseudogene 1 (RPS15AP1) on chromosome
 NG_000952 Homo sapiens ribosomal protein S10-like (RPS10L) pseudogene on chromosome
 NG_000953 Homo sapiens ribosomal protein S23 pseudogene 1 (RPS23P1) on chromosome
 NG_000954 Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3, 12
 NG_000955 Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3, 12
 NG_000956 Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3, 12
 NG_000957 Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3, 12
 NG_000958 Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3, 12
 NG_000959 Homo sapiens similar to DNAJ (HEJ1) pseudogene on chromosome 1
 NG_000960 Homo sapiens ribosomal protein S27a pseudogene 1 (RPS27AP1) on chromosome
 NG_000961 Homo sapiens RNA, U73B small nucleolar (U73B) pseudogene on chromosome
 NG_000962 Homo sapiens ribosomal protein L12 pseudogene 4 (RPL12P4) on chromosome
 NG_000963 Homo sapiens ribosomal protein L38 pseudogene 1 (RPL38P1) on chromosome
 NG_000964 Homo sapiens ribosomal protein L12-like 2 (RPL12L2) pseudogene on chromosome
 NG_000965 Homo sapiens ribosomal protein L27a pseudogene (RPL27AP) on chromosome
 NG_000966 Homo sapiens ribosomal protein L13 pseudogene 2 (RPL13P2) on chromosome
 NG_000967 Homo sapiens ribosomal protein S4-like (RPS4L) pseudogene on chromosome
 NG_000968 Homo sapiens mitochondrial ribosomal protein S16 pseudogene (MRPS16P)
 NG_000969 Homo sapiens ribosomal protein L7a like 2 (RPL7AL2) pseudogene on chromosome

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NG_000970 Homo sapiens ribosomal protein S18 pseudogene 1 (RPS18P1) on chromos
NG_000971 Homo sapiens ribosomal protein L23a pseudogene 6 (RPL23AP6) on chrom
NG_000972 Homo sapiens ribosomal protein S11, pseudogene 1 (RPS11P1) on chromot
NG_000973 Homo sapiens ribosomal protein, large, P0 pseudogene 1 (RPLP0P1) on chr
NG_000974 Homo sapiens ribosomal protein L7a-like 3 (RPL7AL3) pseudogene on chr
NG_000975 Homo sapiens ribosomal protein L15 pseudogene 1 (RPL15P1) on chromos
NG_000977 Homo sapiens mitochondrial ribosomal protein S11 pseudogene 1 (MRPS11
NG_000978 Homo sapiens ribosomal protein L12 pseudogene 3 (RPL12P3) on chromos
NG_000979 Homo sapiens high-mobility group (nonhistone chromosomal) protein 4-like 5
NG_000980 Homo sapiens ribosomal protein L21 pseudogene 4 (RPL21P4) on chromos
NG_000981 Homo sapiens ribosomal protein L7 pseudogene 2 (RPL7P2) on chromosom
NG_000982 Homo sapiens ribosomal protein L37 pseudogene 1 (RPL37P1) on chromos
NG_000983 Homo sapiens ribosomal protein L36 pseudogene 1 (RPL36P1) on chromos
NG_000984 Homo sapiens ribosomal protein S3A pseudogene 3 (RPS3AP3) on chromos
NG_000986 Homo sapiens ribosomal protein L21 pseudogene 2 (RPL21P2) on chromos
NG_000987 Homo sapiens ribosomal protein L35a pseudogene (RPL35AP) on chromos
NG_000988 Homo sapiens ribosomal protein L37a pseudogene 1 (RPL37AP1) on chrom
NG_000989 Homo sapiens ribosomal protein L31 pseudogene 3 (RPL31P3) on chromos
NG_000990 Homo sapiens ribosomal protein L39 pseudogene (RPL39P) on chromosom
NG_000991 Homo sapiens ribosomal protein L17 pseudogene 1 (RPL17P1) on chromos
NG_000994 Homo sapiens ribosomal protein L38 pseudogene 2 (RPL38P2) on chromos
NG_000995 Homo sapiens ribosomal protein S27a pseudogene 2 (RPS27AP2) on chr
NG_000996 Homo sapiens ribosomal protein L31 pseudogene 2 (RPL31P2) on chromos
NG_000997 Homo sapiens ribosomal protein S3 pseudogene 1 (RPS3P1) on chromosom
NG_000999 Homo sapiens ribosomal protein L7A-like 4 (RPL7AL4) pseudogene on chr
NG_001000 Homo sapiens ribosomal protein L24 pseudogene 1 (RPL24P1) on chromos
NG_001001 Homo sapiens ribosomal protein S10 pseudogene 2 (RPS10P2) on chromos
NG_001002 Homo sapiens ribosomal protein L19 pseudogene 1 (RPL19P1) on chromos
NG_001004 Homo sapiens ribosomal protein S3 pseudogene 2 (RPS3P2) on chromosom
NG_001005 Homo sapiens ribosomal protein L7 pseudogene 3 (RPL7P3) on chromosom
NG_001006 Homo sapiens heat shock 10kDa protein 1 (chaperonin 10) pseudogene 1 (H
NG_001007 Homo sapiens cytochrome c oxidase II-like (MTCO2L) pseudogene on chr
NG_001008 Homo sapiens peptidylprolyl isomerase A (cyclophilin A)-like 2 (PPIAL2) pse
NG_001009 Homo sapiens estrogen-related receptor alpha pseudogene (ESRRAP) on c
NG_001010 Homo sapiens similar to RP42 homolog (LOC153893) pseudogene on chr
NG_001012 Homo sapiens TAF2G-like gene (TAF2GL) pseudogene on chromosome 19
NG_001013 Homo sapiens glutaredoxin (thioltransferase) pseudogene (GLRXP) on chr
NG_001014 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) pseudogene 3 (PPI
NG_001016 Homo sapiens C-reactive protein pseudogene (LOC171422) on chromosom
NG_001017 Homo sapiens FUSIP1 pseudogene (pFUSIP1) on chromosome 20
NG_001019 Homo sapiens immunoglobulin heavy locus (IGH.1@) on chromosome 14
NG_001020 Homo sapiens cystatin pseudogene 1 (CSTP1) on chromosome 20
NG_001021 Homo sapiens cytochrome b-5 pseudogene 4 (CYB5P4) on chromosome 20
NG_001022 Homo sapiens RNA binding motif protein, X-linked pseudogene 1 (RBMXP1)
NG_001023 Homo sapiens eukaryotic translation initiation factor 3, subunit 6 48kDa pse
NG_001024 Homo sapiens keratin 18 pseudogene 1 (KRT18P1) on chromosome 6
NG_001025 Homo sapiens peptidylprolyl isomerase (cyclophilin) pseudogene 9 (PPIP9)
NG_001026 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 2 pseudo
NG_001027 Homo sapiens ribosomal protein L23a pseudogene 1 (RPL23AP1) on chrom
NG_001028 Homo sapiens eukaryotic translation termination factor 1 pseudogene 1 (ETF
NG_001029 Homo sapiens makorin, ring finger protein, pseudogene 3 (MKRNP3) on chr
NG_001030 Homo sapiens nuclear fragile X mental retardation protein interacting protein
NG_001031 Homo sapiens ADP-ribosylation factor 4 pseudogene 2 (ARF4P2) on chrom
NG_001032 Homo sapiens cell division cycle 42 pseudogene 1 (CDC42P1) on chromos
NG_001033 Homo sapiens cytochrome c oxidase subunit VIc pseudogene 2 (COX6CP2)
NG_001035 Homo sapiens endosulfine alpha pseudogene (ENSAP) on chromosome 20
NG_001036 Homo sapiens FAT tumor suppressor homolog 1 (Drosophila) pseudogene 1

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NG_001037 Homo sapiens ferritin, light polypeptide pseudogene (FTLP) on chromosome
 NG_001038 Homo sapiens glyceraldehyde-3-phosphate dehydrogenase pseudogene 2 (i
 NG_001040 Homo sapiens glutathione S-transferase M3 pseudogene (GSTM3P) on chro
 NG_001041 Homo sapiens heterogeneous nuclear ribonucleoprotein A1 pseudogene 3 (i
 NG_001043 Homo sapiens keratin 18 pseudogene 3 (KRT18P3) on chromosome 20
 NG_001045 Homo sapiens laminin receptor 1 pseudogene 1 (LAMR1P1) on chromosome
 NG_001046 Homo sapiens proliferation-associated 2G4 pseudogene 2 (PA2G4P2) on ch
 NG_001047 Homo sapiens phosphoglycerate mutase 3, pseudogene (PGAM3P) on chro
 NG_001048 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) pseudogene 2 (PPI
 NG_001049 Homo sapiens peptidylprolyl isomerase (cyclophilin) pseudogene 11 (PPI1'
 NG_001050 Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase,
 NG_001051 Homo sapiens prothymosin, alpha pseudogene 6 (PTMAP6) on chromosome
 NG_001052 Homo sapiens ring finger protein 11B, pseudogene (RNF11B) on chromosom
 NG_001053 Homo sapiens small inducible cytokine subfamily E, member 1 (endothelial n
 NG_001054 Homo sapiens splicing factor 3a, subunit 3 pseudogene (SF3A3P) on chrom
 NG_001055 Homo sapiens synaptosomal-associated protein, 23kDa pseudogene (SNAP
 NG_001056 Homo sapiens small nuclear ribonucleoprotein polypeptide F pseudogene 1
 NG_001057 Homo sapiens spermidine synthase pseudogene 1 (SRMP1) on chromosome
 NG_001058 Homo sapiens suppression of tumorigenicity 13 (colon carcinoma) (Hsp70 in
 NG_001060 Homo sapiens tropomyosin 5, pseudogene (TPM5P) on chromosome 20
 NG_001061 Homo sapiens ubiquitin-conjugating enzyme E2 variant 1 pseudogene 1 (UB
 NG_001062 Homo sapiens exportin, tRNA (nuclear export receptor for tRNAs) pseudoge
 NG_001063 Homo sapiens lysophospholipase I-like (LOC157713) pseudogene on chrom
 NG_001064 Homo sapiens similar to orphan seven transmembrane receptor (RH_II/GuB)
 NG_001065 Homo sapiens RH-II/GuB pseudogene 1 (RH-II/GuBp1) on chromosome 2
 NG_001066 Homo sapiens toll-like receptor 7-like (TLR7-like) pseudogene on chromoso
 NG_001067 Homo sapiens alpha-2-macroglobulin pseudogene (A2MP) on chromosome
 NG_001068 Homo sapiens actin, gamma pseudogene 1 (ACTGP1) on chromosome 3
 NG_001069 Homo sapiens actin, gamma pseudogene 9 (ACTGP9) on chromosome 6
 NG_001070 Homo sapiens adenylate kinase 3 pseudogene 1 (AK3P1) on chromosome 1
 NG_001071 Homo sapiens aldehyde reductase (aldose reductase) pseudogene (ALDRP)
 NG_001073 Homo sapiens S-adenosylmethionine decarboxylase pseudogene 1 (AMDP)
 NG_001074 Homo sapiens v-raf murine sarcoma 3611 viral oncogene homolog pseudoge
 NG_001075 Homo sapiens ADP-ribosylation factor 4 pseudogene (ARF4P) on chromoso
 NG_001076 Homo sapiens ADP-ribosyltransferase 2 pseudogene (RT8 antigen homolog,
 NG_001077 Homo sapiens argininosuccinate synthetase pseudogene 1 (ASSP1) on chro
 NG_001078 Homo sapiens argininosuccinate synthetase pseudogene 3 (ASSP3) on chro
 NG_001080 Homo sapiens activating transcription factor 4 pseudogene (tax-responsive e
 NG_001081 Homo sapiens ATPase, Na⁺/K⁺ transporting, beta polypeptide-like 1 (ATP1E
 NG_001082 Homo sapiens antiqltin-like 1 (ATQL1) pseudogene on chromosome 5
 NG_001083 Homo sapiens antiqltin-like 3 (ATQL3) pseudogene on chromosome 7
 NG_001084 Homo sapiens antiqltin-like 4 (ATQL4) pseudogene on chromosome 10
 NG_001085 Homo sapiens brain cytoplasmic RNA 1, pseudogene 2 (BCYRN1P2) on chr
 NG_001086 Homo sapiens basic transcription factor 3, pseudogene 1 (BTF3P1) on chr
 NG_001087 Homo sapiens solute carrier family 25 (carnitine/acylcarnitine translocase), r
 NG_001088 Homo sapiens calcitonin pseudogene (CALCP) on chromosome 11
 NG_001089 Homo sapiens calmodulin 1 (phosphorylase kinase, delta) pseudogene 1 (C/
 NG_001090 Homo sapiens calmodulin 1 (phosphorylase kinase, delta) pseudogene 2 (C/
 NG_001091 Homo sapiens calmodulin 2 pseudogene 2 (CALM2P2) on chromosome 10
 NG_001092 Homo sapiens cyclin D2 pseudogene (CCND2P) on chromosome 11
 NG_001093 Homo sapiens cyclin D3 pseudogene (CCND3P) on chromosome 6
 NG_001094 Homo sapiens carcinoembryonic antigen-related cell adhesion molecule pse
 NG_001095 Homo sapiens carcinoembryonic antigen-related cell adhesion molecule pse
 NG_001096 Homo sapiens carcinoembryonic antigen-related cell adhesion molecule pse
 NG_001097 Homo sapiens carcinoembryonic antigen-related cell adhesion molecule pse
 NG_001098 Homo sapiens carcinoembryonic antigen-related cell adhesion molecule pse
 NG_001099 Homo sapiens carcinoembryonic antigen-related cell adhesion molecule pse

NG_001100 Homo sapiens carcinoembryonic antigen-related cell adhesion molecule pse
 NG_001101 Homo sapiens carcinoembryonic antigen-related cell adhesion molecule pse
 NG_001102 Homo sapiens carcinoembryonic antigen-related cell adhesion molecule pse
 NG_001103 Homo sapiens carcinoembryonic antigen-related cell adhesion molecule pse
 NG_001104 Homo sapiens carcinoembryonic antigen-related cell adhesion molecule pse
 NG_001105 Homo sapiens cytochrome c oxidase subunit VIa polypeptide 1 pseudogene
 NG_001106 Homo sapiens ceruloplasmin (ferroxidase) pseudogene (CPP) on chromosom
 NG_001107 Homo sapiens crystallin, gamma F pseudogene 1 (CRYGFP1) on chromosom
 NG_001108 Homo sapiens crystallin, gamma G pseudogene 1 (CRYGGP1) on chromosom
 NG_001109 Homo sapiens crystallin, zeta (quinone reductase) pseudogene 1 (CRYZP1)
 NG_001110 Homo sapiens catenin (cadherin-associated protein), alpha pseudogene 1 (C
 NG_001112 Homo sapiens diazepam binding inhibitor-like 2 (pseudogene) (DBIL2) on ch
 NG_001113 Homo sapiens dihydrofolate reductase pseudogene 2 (DHFRP2) on chromo
 NG_001114 Homo sapiens ELK2, member of ETS oncogene family, pseudogene 1 (ELK
 NG_001115 Homo sapiens enolase 1, (alpha) pseudogene (ENO1P) on chromosome 1
 NG_001116 Homo sapiens ferredoxin pseudogene 2 (FDXP2) on chromosome 21
 NG_001117 Homo sapiens ferredoxin pseudogene 1 (FDXP1) on chromosome 20
 NG_001118 Homo sapiens ferrochelatase pseudogene (FECHP) on chromosome 3
 NG_001119 Homo sapiens forkhead box O3B (FOXO3B) pseudogene on chromosome 1
 NG_001120 Homo sapiens forkhead box O1B (FOXO1B) pseudogene on chromosome 5
 NG_001121 Homo sapiens ferritin, heavy polypeptide pseudogene 2 (FTHP2) on chromo
 NG_001122 Homo sapiens fucosidase, alpha-L- 1, tissue pseudogene (FUCA1P) on chro
 NG_001123 Homo sapiens glyceraldehyde-3-phosphate dehydrogenase pseudogene 1 (G
 NG_001124 Homo sapiens GDP dissociation inhibitor 2 pseudogene (GDI2P) on chromo
 NG_001125 Homo sapiens glycoprotein, alpha-galactosyltransferase 1 (GGTA1) pseudog
 NG_001126 Homo sapiens glycerol kinase pseudogene 1 (GKP1) on chromosome 5
 NG_001127 Homo sapiens glycerol kinase pseudogene 6 (GKP6) on chromosome X
 NG_001128 Homo sapiens glutamate dehydrogenase pseudogene 2 (GLUDP2) on chro
 NG_001129 Homo sapiens glutamate dehydrogenase pseudogene 5 (GLUDP5) on chro
 NG_001130 Homo sapiens GM2 ganglioside activator pseudogene (GM2AP) on chromos
 NG_001131 Homo sapiens G protein-coupled receptor 32, pseudogene (GPR32P) on chr
 NG_001132 Homo sapiens G protein-coupled receptor 33, pseudogene (GPR33) on chr
 NG_001133 Homo sapiens G protein-coupled receptor kinase 6 pseudogene (GRK6PS) c
 NG_001134 Homo sapiens glutathione peroxidase pseudogene 2 (GPXP2) on chromosom
 NG_001135 Homo sapiens glutathione S-transferase A pseudogene 1 (GSTAP1) on chro
 NG_001136 Homo sapiens gulonolactone (L-) oxidase pseudogene (GULOPO) on chromo
 NG_001141 Homo sapiens high-mobility group (nonhistone chromosomal) protein 17 pse
 NG_001142 Homo sapiens high-mobility group (nonhistone chromosomal) protein 17 pse
 NG_001144 Homo sapiens heat shock 70kDa protein pseudogene 1 (HSPAP1) on chrom
 NG_001145 Homo sapiens heat shock 60kDa protein 1 (chaperonin) pseudogene 1 (HSF
 NG_001146 Homo sapiens heat shock 60kDa protein 1 (chaperonin) pseudogene 2 (HSF
 NG_001147 Homo sapiens heat shock 60kDa protein 1 (chaperonin) pseudogene 3 (HSF
 NG_001148 Homo sapiens heat shock 60kDa protein 1 (chaperonin) pseudogene 4 (HSF
 NG_001149 Homo sapiens iduronate 2-sulfatase pseudogene 1 (IDSP1) on chromosome
 NG_001150 Homo sapiens interferon, omega 15 (pseudogene) (IFNWP15) on chromosom
 NG_001151 Homo sapiens recombining binding protein suppressor of hairless (Drosophil
 NG_001152 Homo sapiens IMP (inosine monophosphate) dehydrogenase-like 1 (IMPDH
 NG_001153 Homo sapiens Kallmann syndrome sequence pseudogene (KALP) on chromo
 NG_001155 Homo sapiens lactate dehydrogenase B pseudogene (LDHBP) on chromosom
 NG_001156 Homo sapiens melanoma antigen, family A, 7, pseudogene (MAGEA7) on ch
 NG_001157 Homo sapiens MRE11 meiotic recombination 11 homolog B (S. cerevisiae) (I
 NG_001158 Homo sapiens metallothionein 2 pseudogene 1 (processed) (MT2P1) on chr
 NG_001159 Homo sapiens methyltetrahydrofolate dehydrogenase (NADP)- depende
 NG_001160 Homo sapiens metaxin 1 pseudogene (MTX1P) on chromosome 1
 NG_001161 Homo sapiens NADH dehydrogenase (ubiquinone) flavoprotein 2 pseudog
 NG_001162 Homo sapiens nucleophosmin 1 (nucleolar phosphoprotein B23, numatrin) p
 NG_001163 Homo sapiens nucleophosmin 1 (nucleolar phosphoprotein B23, numatrin) p

NG_001164 Homo sapiens nucleophosmin 1 (nucleolar phosphoprotein B23, numatrin) p
 NG_001165 Homo sapiens nucleophosmin 1 (nucleolar phosphoprotein B23, numatrin) p
 NG_001166 Homo sapiens nucleophosmin 1 (nucleolar phosphoprotein B23, numatrin) p
 NG_001167 Homo sapiens nucleophosmin 1 (nucleolar phosphoprotein B23, numatrin) p
 NG_001169 Homo sapiens ornithine aminotransferase-like 3 (pseudogene) (OATL3) on c
 NG_001170 Homo sapiens phosphoglycerate kinase 1, pseudogene 1 (PGK1P1) on chro
 NG_001171 Homo sapiens prohibitin pseudogene 1 (PHBP1) on chromosome 6
 NG_001172 Homo sapiens phosphorylase kinase, beta pseudogene 1 (PHKBP1) on chro
 NG_001173 Homo sapiens phosphorylase kinase, beta pseudogene 2 (PHKBP2) on chro
 NG_001174 Homo sapiens phosphatidylinositol glycan, class A, pseudogene 1 (PIGAP1)
 NG_001175 Homo sapiens phosphatidylinositol glycan, class C, pseudogene 1 (PIGCP1)
 NG_001176 Homo sapiens phosphatidylinositol glycan, class F, pseudogene 1 (PIGFP1)
 NG_001177 Homo sapiens peptidylprolyl isomerase (cyclophilin) pseudogene 1 (PPIP1) c
 NG_001178 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 8 pseudo
 NG_001179 Homo sapiens protein S pseudogene (beta) (PROSP) on chromosome 3
 NG_001180 Homo sapiens prothymosin, alpha pseudogene 3 (gene sequence 34) (PTM/
 NG_001181 Homo sapiens prothymosin, alpha pseudogene 4 (gene sequence 112) (PTM/
 NG_001182 Homo sapiens protein tyrosine phosphatase type IVA pseudogene 2 (PTP4A
 NG_001183 Homo sapiens RNA, U1 small nuclear pseudogene 1 (RNU1P1) on chromos
 NG_001184 Homo sapiens RNA, U1 small nuclear pseudogene 2 (RNU1P2) on chromos
 NG_001185 Homo sapiens RNA, U3 small nuclear pseudogene 1 (RNU3P1)
 NG_001186 Homo sapiens RNA, U4 small nuclear pseudogene 1 (U4/7) (RNU4P1) on ch
 NG_001187 Homo sapiens RNA, U4 small nuclear pseudogene 2 (U4/14) (RNU4P2) on c
 NG_001188 Homo sapiens RNA, U7 small nuclear pseudogene 1 (RNU7P1) on chromos
 NG_001189 Homo sapiens RNA, U7 small nuclear pseudogene 2 (RNU7P2) on chromos
 NG_001190 Homo sapiens RNA, U7 small nuclear pseudogene 3 (RNU7P3) on chromos
 NG_001191 Homo sapiens RNA, U7 small nuclear pseudogene 4 (RNU7P4) on chromos
 NG_001192 Homo sapiens ribosomal protein L9 pseudogene 1 (RPL9P1) on chromosom
 NG_001193 Homo sapiens ribosomal protein L7 pseudogene (RPL7P) on chromosome 5
 NG_001194 Homo sapiens sterol-C4-methyl oxidase pseudogene (SC4MOP) on chromo
 NG_001195 Homo sapiens SHC (Src homology 2 domain containing) transforming protei
 NG_001196 Homo sapiens steroid-5-alpha-reductase, alpha polypeptide pseudogene 1 (;
 NG_001197 Homo sapiens steroid sulfatase (microsomal) pseudogene (STSP) on chrom
 NG_001198 Homo sapiens eukaryotic translation termination factor 1 pseudogene 2 (ETF
 NG_001199 Homo sapiens TAR (HIV) RNA binding protein 2 pseudogene (TARBP2P) on
 NG_001200 Homo sapiens transcription elongation factor A (SII), 1 pseudogene (TCEA11
 NG_001201 Homo sapiens thioredoxin-dependent peroxide reductase 2 (thiol-specific an
 NG_001202 Homo sapiens transcription factor Dp-1 pseudogene (TFDP1P) on chromoso
 NG_001204 Homo sapiens tRNA leucine (AAG) pseudogene 1 (TRLP1) on chromosome
 NG_001205 Homo sapiens tRNA methionine elongator pseudogene 1 (TRMEP1) on chro
 NG_001206 Homo sapiens tubulin, beta polypeptide pseudogene 1 (TUBBP1) on chro
 NG_001207 Homo sapiens ubiquitin A-52 residue ribosomal protein fusion product 1 pse
 NG_001208 Homo sapiens ubiquitin A-52 residue ribosomal protein fusion product 1 pse
 NG_001209 Homo sapiens ubiquitin-conjugating enzyme E2L 4 (UBE2L4) pseudogene o
 NG_001210 Homo sapiens ubiquitin protein ligase E3A pseudogene 2 (UBE3AP2) on chr
 NG_001211 Homo sapiens urate oxidase (UOX) pseudogene on chromosome 1
 NG_001212 Homo sapiens von Willebrand factor pseudogene (VWF) on chromosome 2
 NG_001213 Homo sapiens v-yes-1 Yamaguchi sarcoma viral oncogene homolog pseudo
 NG_001214 Homo sapiens zinc finger protein 75b (ZNF75B) pseudogene on chromosom
 NG_001215 Homo sapiens interferon-induced protein with tetraatricopeptide repeats 1, psi
 NG_001216 Homo sapiens a disintegrin and metalloproteinase domain 1 (fertilin alpha) p
 NG_001217 Homo sapiens G protein-coupled receptor 53, pseudogene (GPR53P) on chr
 NG_001218 Homo sapiens synaptogyrin 2 pseudogene (SYNGR2P) on chromosome 15
 NG_001219 Homo sapiens cytochrome c oxidase subunit VIc pseudogene 1 (COX6CP1)
 NG_001220 Homo sapiens cytochrome c oxidase subunit Va pseudogene 1 (COX5AP1)
 NG_001221 Homo sapiens cytochrome c oxidase subunit VIc pseudogene 1 (COX7CP1)
 NG_001222 Homo sapiens v-myc myelocytomatosis viral oncogene homolog 3 pseudoge

NG_001223 Homo sapiens voltage-dependent anion channel 1-like pseudogene (VDAC1
NG_001224 Homo sapiens voltage-dependent anion channel 1 pseudogene (VDAC1P) o
NG_001226 Homo sapiens H2B histone family, member O (H2BFO) pseudogene
NG_001228 Homo sapiens cold shock domain protein A pseudogene 1 (CSDAP1) on chr
NG_001229 Homo sapiens nuclear distribution gene C homolog (A. nidulans) pseudogen
NG_001230 Homo sapiens nuclear distribution gene C homolog (A. nidulans) pseudogen
NG_001231 Homo sapiens nucleophosmin 1 (nucleolar phosphoprotein B23, numatrin) p
NG_001232 Homo sapiens nucleophosmin 1 (nucleolar phosphoprotein B23, numatrin) p
NG_001233 Homo sapiens nucleophosmin 1 (nucleolar phosphoprotein B23, numatrin) p
NG_001234 Homo sapiens macrophage stimulating, pseudogene 8 (MSTP8) on chromos
NG_001235 Homo sapiens macrophage stimulating, pseudogene 7 (MSTP7) on chromos
NG_001236 Homo sapiens macrophage stimulating, pseudogene 6 (MSTP6) on chromos
NG_001237 Homo sapiens macrophage stimulating, pseudogene 5 (MSTP5) on chromos
NG_001238 Homo sapiens macrophage stimulating, pseudogene 4 (MSTP4) on chromos
NG_001239 Homo sapiens macrophage stimulating, pseudogene 3 (MSTP3) on chromos
NG_001240 Homo sapiens macrophage stimulating, pseudogene 2 (MSTP2) on chromos
NG_001241 Homo sapiens macrophage stimulating, pseudogene 1 (MSTP1) on chromos
NG_001242 Homo sapiens teratocarcinoma-derived growth factor 5, pseudogene (TDGF1
NG_001243 Homo sapiens teratocarcinoma-derived growth factor 4, pseudogene (TDGF1
NG_001244 Homo sapiens teratocarcinoma-derived growth factor 2, pseudogene (TDGF1
NG_001245 Homo sapiens glycosylphosphatidylinositol anchor attachment 1 pseudogen
NG_001246 Homo sapiens ATP-binding cassette, sub-family D (ALD), member 1, pseud
NG_001247 Homo sapiens protocadherin alpha 14 pseudogene (PCDHA14) on chromos
NG_001248 Homo sapiens RNA, U6 small nuclear pseudogene 1 (RNU6P1) on chromos
NG_001249 Homo sapiens RNA, U4 small nuclear pseudogene 8 (RNU4P8) on chromos
NG_001250 Homo sapiens RNA, U4 small nuclear pseudogene 5 (RNU4P5) on chromos
NG_001251 Homo sapiens RNA, U4 small nuclear pseudogene 4 (RNU4P4) on chromos
NG_001252 Homo sapiens RNA, U4 small nuclear pseudogene 3 (RNU4P3) on chromos
NG_001253 Homo sapiens RNA, U3 small nucleolar pseudogene 4 (RNU3P4) on chrom
NG_001254 Homo sapiens RNA, U3 small nucleolar pseudogene 3 (RNU3P3) on chrom
NG_001255 Homo sapiens RNA, U3 small nucleolar pseudogene 2 (RNU3P2) on chrom
NG_001256 Homo sapiens RNA, U2 small nuclear pseudogene 3 (RNU2P3) on chromos
NG_001257 Homo sapiens RNA, U2 small nuclear pseudogene 2 (RNU2P2) on chromos
NG_001258 Homo sapiens RNA, U2 small nuclear pseudogene 1 (RNU2P1)
NG_001259 Homo sapiens RNA, U1 small nuclear pseudogene 10 (RNU1P10) on chrom
NG_001260 Homo sapiens RNA, U1 small nuclear pseudogene 9 (RNU1P9) on chromos
NG_001261 Homo sapiens RNA, U1 small nuclear pseudogene 8 (RNU1P8) on chromos
NG_001262 Homo sapiens RNA, U1 small nuclear pseudogene 7 (RNU1P7)
NG_001263 Homo sapiens RNA, U1 small nuclear pseudogene 6 (RNU1P6) on chromos
NG_001264 Homo sapiens RNA, U1 small nuclear pseudogene 5 (RNU1P5) on chromos
NG_001265 Homo sapiens ATP-binding cassette, sub-family D (ALD), member 1, pseud
NG_001266 Homo sapiens poly(A) binding protein, cytoplasmic, pseudogene 3 (PABPCF
NG_001267 Homo sapiens poly(A) binding protein, cytoplasmic, pseudogene 2 (PABPCF
NG_001268 Homo sapiens poly(A) binding protein, cytoplasmic, pseudogene 1 (PABPCF
NG_001269 Homo sapiens ATP-binding cassette, sub-family D (ALD), member 1, pseud
NG_001270 Homo sapiens ATP-binding cassette, sub-family D (ALD), member 1, pseud
NG_001271 Homo sapiens molybdenum cofactor synthesis 1 pseudogene 1 (MOCS1P1)
NG_001272 Homo sapiens G protein-coupled receptor 79 pseudogene (GPR79) on chr
NG_001273 Homo sapiens mitogen-activated protein kinase kinase 1 pseudogene 1 (MA
NG_001274 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 14B pse
NG_001275 Homo sapiens chemokine (C-X-C motif) ligand 1 pseudogene (CXCL1P) on
NG_001276 Homo sapiens nuclease sensitive element binding protein 1 pseudogene (NE
NG_001277 Homo sapiens capping protein (actin filament) muscle Z-line, alpha 1 pseud
NG_001278 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B28 pseudoge
NG_001279 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B27 pseudoge
NG_001280 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B26 pseudoge
NG_001281 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B25 pseudoge

NG_001282 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B24 pseudogene
 NG_001286 Homo sapiens fatty acid binding protein 3, pseudogene 2 (FABP3P2) on chr
 NG_001287 Homo sapiens uracil-DNA glycosylase pseudogene 1 (UNGFP1) on chromos
 NG_001288 Homo sapiens uracil-DNA glycosylase pseudogene 2 (UNGFP2) on chromos
 NG_001289 Homo sapiens Zn-15 related zinc finger protein RLF pseudogene (RLFP)
 NG_001290 Homo sapiens 6-pyruvoyltetrahydropterin synthase pseudogene (PTS-P1) or
 NG_001291 Homo sapiens recombining binding protein suppressor of hairless (Drosophil
 NG_001292 Homo sapiens recombining binding protein suppressor of hairless (Drosophil
 NG_001293 Homo sapiens COX17 pseudogene (LOC81993) on chromosome 13
 NG_001294 Homo sapiens cytochrome P450, subfamily 51 pseudogene 1 (CYP51P1) on
 NG_001295 Homo sapiens ribosomal protein S19 pseudogene 1 (RPS19P1) on chromos
 NG_001296 Homo sapiens ribosomal protein S19 pseudogene 2 (RPS19P2) on chromos
 NG_001297 Homo sapiens thioredoxin 1 pseudogene 2 (LOC93202) on chromosome 10
 NG_001298 Homo sapiens family with sequence similarity 8, member A5 pseudogene (F
 NG_001299 Homo sapiens family with sequence similarity 8, member A6 pseudogene (F
 NG_001300 Homo sapiens cytochrome c oxidase subunit VIIb pseudogene 1 (COX7BP1
 NG_001301 Homo sapiens ribosomal protein L37 pseudogene 2 (RPL37P2) on chromos
 NG_001302 Homo sapiens ribosomal protein L3 pseudogene 2 (RPL3P2) on chromosom
 NG_001303 Homo sapiens thioredoxin 1 pseudogene 4 (LOC124974) on chromosome 1
 NG_001305 Homo sapiens mitogen-activated protein kinase kinase 4 pseudogene (LOC1
 NG_001306 Homo sapiens FBR-MuSV-associated ubiquitously expressed (fox derived) p
 NG_001307 Homo sapiens protein phosphatase 2, regulatory subunit B (B56), gamma iso
 NG_001308 Homo sapiens hydroxysteroid (17-beta) dehydrogenase 7 pseudogene 1 (HE
 NG_001309 Homo sapiens thioredoxin 1 pseudogene 1 (LOC151276) on chromosome 2
 NG_001311 Homo sapiens deleted in split-hand/split-foot 1 pseudogene (DSS1P1) on chr
 NG_001313 Homo sapiens lactate dehydrogenase pseudogene (LOC158222) on chromos
 NG_001314 Homo sapiens PHD finger protein 10 pseudogene 1 (PHF10P1) on chromos
 NG_001315 Homo sapiens chromobox homolog 3 gamma pseudogene (LOC159770) on
 NG_001316 Homo sapiens ribosomal protein L7a pseudogene 2 (RPL7AP2) on chromos
 NG_001317 Homo sapiens ribosomal protein L7a pseudogene 3 (RPL7AP3) on chromos
 NG_001318 Homo sapiens HSP40 pseudogene (HSP40) on chromosome 2
 NG_001319 Homo sapiens ubiquitin-conjugating enzyme-like (UBCH7N2) pseudogene on
 NG_001321 Homo sapiens thioredoxin 1 pseudogene 6 (AF357533) on chromosome 1
 NG_001322 Homo sapiens thioredoxin 1 pseudogene 7 (AF357534) on chromosome 2
 NG_001323 Homo sapiens glutaredoxin pseudogene 2 (GLRXP2) on chromosome 14
 NG_001324 Homo sapiens thioredoxin 1 pseudogene 5 (AF357532) on chromosome 1
 NG_001325 Homo sapiens EPF5 pseudogene (EPF5) on chromosome 9
 NG_001326 Homo sapiens EPF8 pseudogene (EPF8) on chromosome 16
 NG_001328 Homo sapiens sperm autoantigenic protein 17 pseudogene 1 (SPA17P1) on
 NG_001329 Homo sapiens high mobility group AT-hook 1-like 2 (HMGA1L2) pseudogene
 NG_001330 Homo sapiens thioredoxin 1 pseudogene 3 (AF357530) on chromosome 4
 NG_001331 Homo sapiens 3-oxoacid CoA transferase 2 pseudogene (OXC2P2) on chro
 NG_001332 Homo sapiens T cell receptor alpha delta locus (TCRA/TCRD) on chromosom
 NG_001333 Homo sapiens T cell receptor beta locus (TRB@) on chromosome 7
 NG_001334 Homo sapiens growth hormone locus (GH@) on chromosome 17
 NG_001335 Homo sapiens genomic large histone family cluster (HFL@) on chromosome
 NG_001336 Homo sapiens T cell receptor gamma locus (TRG@) on chromosome 7
 NG_001337 Homo sapiens T cell receptor beta variable orphans on chromosome 9 (TRB
 NG_001526 Homo sapiens a disintegrin and metalloproteinase domain 3b (cytistatin 2) (F
 NG_001528 Homo sapiens phosphoglycerate kinase 1, pseudogene 2 (PGK1P2) on chro
 NG_001529 Homo sapiens tyrosinase-like (TYRL) pseudogene on chromosome 11
 NG_001531 Homo sapiens similar to tyrosine 3-monooxygenase/tryptophan 5-monooxyge
 NG_001532 Homo sapiens similar to casein kinase 1, alpha 1 (LOC120321) pseudogene
 NG_001533 Homo sapiens proteasome (prosome, macropain) subunit, beta type, 3 pseu
 NG_001534 Homo sapiens legumain 2 pseudogene (LGMN2P) on chromosome 13
 NG_001535 Homo sapiens brain creatine kinase pseudogene (LOC124144) on chromos
 NG_001537 Homo sapiens coactosin-like, Smith Magenis syndrome chromosome region

NG_001538 Homo sapiens proteasome (prosome, macropain) subunit, beta type, 3 pseud
 NG_001539 Homo sapiens mitochondrial ribosomal protein L9 pseudogene (MRPL9P1) c
 NG_001540 Homo sapiens ATP synthase, H⁺ transporting, mitochondrial F0 complex, su
 NG_001541 Homo sapiens synaptogyrin 2 pseudogene (LOC138916) on chromosome 9
 NG_001542 Homo sapiens similar to plasmolipin (LOC139061) pseudogene on chromos
 NG_001543 Homo sapiens peptidyl prolyl isomerase H (cyclophilin H) pseudogene (LOC
 NG_001544 Homo sapiens eukaryotic translation initiation factor 3, subunit 6 interacting f
 NG_001545 Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase acti
 NG_001546 Homo sapiens similar to yeast Upf3, variant A pseudogene (LOC147150) on
 NG_001547 Homo sapiens lectin, galactoside-binding, soluble, 9 (galectin 9) pseudogene
 NG_001548 Homo sapiens similar to yeast Upf3, variant A pseudogene (LOC147226) on
 NG_001549 Homo sapiens son-pseudogene (LOC148300) on chromosome 1
 NG_001550 Homo sapiens u2 small nuclear ribonucleoprotein A' pseudogene (LOC1504:
 NG_001551 Homo sapiens u2 small nuclear ribonucleoprotein polypeptide A' pseudogeni
 NG_001552 Homo sapiens RAB6C, member RAS oncogene family pseudogene (LOC15(
 NG_001553 Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase acti
 NG_001554 Homo sapiens RAB6C, member RAS oncogene family pseudogene (LOC15(
 NG_001555 Homo sapiens peptidylprolyl isomerase (cyclophilin)-like 1 pseudogene 1 (Pf
 NG_001556 Homo sapiens protease (prosome, macropain) 26S subunit, ATPase, 1 pseu
 NG_001557 Homo sapiens similar to ART-4 protein (LOC152594) pseudogene on chromi
 NG_001558 Homo sapiens signal recognition particle 72kD pseudogene (LOC153932) or
 NG_001559 Homo sapiens protein phosphatase 2, regulatory subunit B (B56), epsilon lac
 NG_001560 Homo sapiens cell division cycle 20 pseudogene (LOC157956) on chromoso
 NG_001561 Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase acti
 NG_001562 Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase acti
 NG_001563 Homo sapiens protein kinase C, iota pseudogene (LOC158948) on chromosi
 NG_001564 Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 6 p
 NG_001566 Homo sapiens peptidylprolyl isomerase A (cyclophilin A)-like (PPIAL) pseud
 NG_001567 Homo sapiens ribonuclease H1 pseudogene 3 (RNASEH1P3) on chromosom
 NG_001568 Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) pseudogene
 NG_001569 Homo sapiens similar to high mobility group protein-R (LOC201958) pseudog
 NG_001571 Homo sapiens ribosomal protein L36a pseudogene (dJ507115.1) on chromo
 NG_001572 Homo sapiens cyclin-dependent kinase 7 pseudogene (CDK7PS) on chromo
 NG_001573 Homo sapiens TAF13 RNA polymerase II, TATA box binding protein (TBP)-a
 NG_001574 Homo sapiens TBP-associated factor 9-like pseudogene (LOC246135) on ch
 NG_001576 Homo sapiens deltaNEMO (deltaNEMO) pseudogene on chromosome X
 NG_001577 Homo sapiens CDC28 protein kinase regulatory subunit 1A (CKS1A) pseudo
 NG_001578 Homo sapiens CDC28 protein kinase regulatory subunit 1B pseudogene 2 (C
 NG_001579 Homo sapiens CDC28 protein kinase regulatory subunit 1B pseudogene 3 (C
 NG_001580 Homo sapiens YWHAQ pseudogene 2 (YWHAQF2) on chromosome 22
 NG_001581 Homo sapiens transcription elongation factor B (SII), polypeptide 2 (18kD, el
 NG_001582 Homo sapiens transcription elongation factor B (SII), polypeptide 2 (18kD, el
 NG_001583 Homo sapiens ubiquitin carrier protein E2-EPF pseudogene (LOC246719) or
 NG_001584 Homo sapiens dynein, cytoplasmic, light polypeptide pseudogene (LOC2467
 NG_001585 Homo sapiens ATP synthase, H⁺ transporting, mitochondrial F0 complex, su
 NG_001586 Homo sapiens peptidyl prolyl isomerase H (cyclophilin H) pseudogene 1 (PP
 NG_001587 Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase acti
 NG_001588 Homo sapiens homolog of C. elegans smu-1 pseudogene (LOC246784) on c
 NG_001589 Homo sapiens PTD004 pseudogene (LOC246785) on chromosome 22
 NG_001590 Homo sapiens signal recognition particle 68kD pseudogene (LOC252840) or
 NG_001591 Homo sapiens signal recognition particle 68kD pseudogene (LOC252841) or
 NG_001592 Homo sapiens karyopherin (importin) beta 2 pseudogene (LOC252966) on cl
 NG_002151 Homo sapiens olfactory receptor, family 1, subfamily E, member 3 pseudoge
 NG_002153 Homo sapiens olfactory receptor, family 1, subfamily P, member 1 pseudoge
 NG_002154 Homo sapiens olfactory receptor, family 8, subfamily C, member 1 pseudoge
 NG_002156 Homo sapiens olfactory receptor, family 8, subfamily B, member 1 pseudoge
 NG_002158 Homo sapiens olfactory receptor, family 7, subfamily E, member 41 pseudog

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NG_002160 Homo sapiens olfactory receptor, family 7, subfamily E, member 5 pseudog
NG_002162 Homo sapiens olfactory receptor, family 7, subfamily E, member 4 pseudog
NG_002164 Homo sapiens olfactory receptor, family 7, subfamily E, member 87 pseudog
NG_002166 Homo sapiens olfactory receptor, family 7, subfamily E, member 2 pseudog
NG_002168 Homo sapiens olfactory receptor, family 7, subfamily E, member 15 pseudog
NG_002170 Homo sapiens olfactory receptor, family 7, subfamily A, member 3 pseudog
NG_002172 Homo sapiens olfactory receptor, family 5, subfamily D, member 3 pseudog
NG_002173 Homo sapiens olfactory receptor, family 4, subfamily A, member 1 pseudog
NG_002175 Homo sapiens olfactory receptor, family 7, subfamily E, member 14 pseudog
NG_002177 Homo sapiens olfactory receptor, family 7, subfamily E, member 13 pseudog
NG_002179 Homo sapiens olfactory receptor, family 7, subfamily E, member 12 pseudog
NG_002181 Homo sapiens olfactory receptor, family 7, subfamily E, member 11 pseudog
NG_002183 Homo sapiens olfactory receptor, family 7, subfamily E, member 10 pseudog
NG_002184 Homo sapiens olfactory receptor, family 4, subfamily C, member 1 pseudog
NG_002185 Homo sapiens olfactory receptor, family 4, subfamily H, member 8 pseudog
NG_002189 Homo sapiens olfactory receptor, family 7, subfamily E, member 16 pseudog
NG_002191 Homo sapiens olfactory receptor, family 7, subfamily E, member 43 pseudog
NG_002195 Homo sapiens olfactory receptor, family 9, subfamily A, member 1, pseudog
NG_002196 Homo sapiens olfactory receptor, family 12, subfamily D, member 1 pseudog
NG_002199 Homo sapiens olfactory receptor, family 51, subfamily A, member 1 pseudog
NG_002200 Homo sapiens olfactory receptor, family 10, subfamily G, member 1 pseudog
NG_002201 Homo sapiens olfactory receptor, family 10, subfamily B, member 1 pseudog
NG_002202 Homo sapiens olfactory receptor, family 8, subfamily B, member 7 pseudog
NG_002203 Homo sapiens olfactory receptor, family 8, subfamily B, member 6 pseudog
NG_002204 Homo sapiens olfactory receptor, family 8, subfamily B, member 5 pseudog
NG_002205 Homo sapiens olfactory receptor, family 7, subfamily H, member 1 pseudog
NG_002207 Homo sapiens olfactory receptor, family 7, subfamily E, member 8 pseudog
NG_002211 Homo sapiens olfactory receptor, family 7, subfamily A, member 33 pseudog
NG_002212 Homo sapiens olfactory receptor, family 7, subfamily D, member 1 pseudog
NG_002215 Homo sapiens olfactory receptor, family 7, subfamily A, member 11 pseudog
NG_002216 Homo sapiens olfactory receptor, family 2, subfamily N, member 1 pseudog
NG_002218 Homo sapiens vomeronasal 1 receptor 7 pseudog (VN1R7P) on chromo
NG_002219 Homo sapiens olfactory receptor, family 56, subfamily A, member 5 (OR56A)
NG_002220 Homo sapiens olfactory receptor, family 52, subfamily X, member 1 pseudog
NG_002221 Homo sapiens olfactory receptor, family 7, subfamily E, member 94 pseudog
NG_002225 Homo sapiens olfactory receptor, family 7, subfamily E, member 93 pseudog
NG_002229 Homo sapiens olfactory receptor, family 4, subfamily C, member 7 pseudog
NG_002232 Homo sapiens olfactory receptor, family 51, subfamily P, member 1 pseudog
NG_002233 Homo sapiens olfactory receptor, family 9, subfamily L, member 1 pseudog
NG_002234 Homo sapiens olfactory receptor, family 52, subfamily J, member 1 pseudog
NG_002235 Homo sapiens olfactory receptor, family 4, subfamily K, member 7 pseudog
NG_002236 Homo sapiens olfactory receptor, family 4, subfamily P, member 1 pseudog
NG_002238 Homo sapiens olfactory receptor, family 1, subfamily AA, member 1 pseudog
NG_002239 Homo sapiens olfactory receptor, family 2, subfamily AD, member 1 pseudog
NG_002240 Homo sapiens olfactory receptor, family 4, subfamily K, member 6 pseudog
NG_002241 Homo sapiens olfactory receptor, family 7, subfamily E, member 91 pseudog
NG_002242 Homo sapiens olfactory receptor, family 4, subfamily K, member 4 pseudog
NG_002243 Homo sapiens olfactory receptor, family 4, subfamily N, member 1 pseudog
NG_002246 Homo sapiens olfactory receptor, family 13, subfamily C, member 1 pseudog
NG_002247 Homo sapiens olfactory receptor, family 4, subfamily C, member 5 (OR4C5)
NG_002250 Homo sapiens olfactory receptor, family 4, subfamily C, member 2 pseudog
NG_002251 Homo sapiens olfactory receptor, family 7, subfamily E, member 83 pseudog
NG_002252 Homo sapiens olfactory receptor, family 51, subfamily J, member 1 (OR51J1
NG_002253 Homo sapiens olfactory receptor, family 5, subfamily BD, member 1 pseudog
NG_002254 Homo sapiens olfactory receptor, family 10, subfamily D, member 5 pseudog
NG_002255 Homo sapiens olfactory receptor, family 10, subfamily G, member 6 (OR10G
NG_002256 Homo sapiens olfactory receptor, family 7, subfamily E, member 97 pseudog

NG_002257 Homo sapiens olfactory receptor, family 2, subfamily AF, member 1 pseudog
 NG_002258 Homo sapiens olfactory receptor, family 7, subfamily L, member 1 pseudog
 NG_002259 Homo sapiens olfactory receptor, family 10, subfamily R, member 1 pseudog
 NG_002260 Homo sapiens olfactory receptor, family 10, subfamily T, member 1 pseudog
 NG_002261 Homo sapiens olfactory receptor, family 10, subfamily G, member 5 pseudog
 NG_002262 Homo sapiens olfactory receptor, family 52, subfamily S, member 1 pseudog
 NG_002263 Homo sapiens olfactory receptor, family 51, subfamily A, member 5 pseudog
 NG_002264 Homo sapiens olfactory receptor, family 4, subfamily C, member 10 pseudog
 NG_002265 Homo sapiens olfactory receptor, family 4, subfamily R, member 1 pseudog
 NG_002266 Homo sapiens olfactory receptor, family 52, subfamily J, member 2 pseudog
 NG_002267 Homo sapiens olfactory receptor, family 4, subfamily C, member 9 pseudog
 NG_002268 Homo sapiens olfactory receptor, family 51, subfamily A, member 3 pseudog
 NG_002269 Homo sapiens olfactory receptor, family 52, subfamily E, member 3 pseudog
 NG_002270 Homo sapiens olfactory receptor, family 4, subfamily V, member 1 pseudog
 NG_002271 Homo sapiens olfactory receptor, family 7, subfamily E, member 90 pseudog
 NG_002272 Homo sapiens olfactory receptor, family 7, subfamily E, member 89 pseudog
 NG_002273 Homo sapiens olfactory receptor, family 2, subfamily AL, member 1 pseudog
 NG_002274 Homo sapiens olfactory receptor, family 6, subfamily J, member 1 (OR6J1) p
 NG_002275 Homo sapiens olfactory receptor, family 4, subfamily C, member 2 pseudog
 NG_002277 Homo sapiens olfactory receptor, family 5, subfamily D, member 2 pseudog
 NG_002278 Homo sapiens olfactory receptor, family 7, subfamily A, member 18 pseudog
 NG_002279 Homo sapiens olfactory receptor, family 9, subfamily I, member 2 pseudog
 NG_002281 Homo sapiens olfactory receptor, family 5, subfamily M, member 13 pseudog
 NG_002282 Homo sapiens olfactory receptor, family 6, subfamily L, member 1 pseudog
 NG_002284 Homo sapiens ubiquitin B pseudogene 1 (UBBP1) on chromosome 2
 NG_002285 Homo sapiens ubiquitin B pseudogene 4 (UBBP4) on chromosome 17
 NG_002286 Homo sapiens ubiquitin B pseudogene 3 (UBBP3) on chromosome 2
 NG_002287 Homo sapiens ubiquitin B pseudogene 2 (UBBP2) on chromosome 1
 NG_002288 Homo sapiens suppressor of btmD6 homolog pseudogene (LOC122145) on
 NG_002289 Homo sapiens serine hydroxymethyltransferase 1 (soluble) pseudogene (SH
 NG_002290 Homo sapiens DKFZP434J193-like pseudogene (LOC258308) on chromos
 NG_002291 Homo sapiens splicing factor 3b, subunit 4, 49kD pseudogene (LOC260329)
 NG_002298 Homo sapiens olfactory receptor, family 1, subfamily D, member 3 pseudog
 NG_002299 Homo sapiens olfactory receptor, family 5, subfamily G, member 1 pseudog
 NG_002302 Homo sapiens olfactory receptor, family 1, subfamily R, member 1 pseudog
 NG_002305 Homo sapiens olfactory receptor, family 5, subfamily B, member 1 pseudog
 NG_002306 Homo sapiens olfactory receptor, family 5, subfamily B, member 10 pseudog
 NG_002307 Homo sapiens olfactory receptor, family 4, subfamily H, member 6 pseudog
 NG_002311 Homo sapiens olfactory receptor, family 7, subfamily A, member 15 pseudog
 NG_002315 Homo sapiens olfactory receptor, family 7, subfamily E, member 26 pseudog
 NG_002316 Homo sapiens olfactory receptor, family 7, subfamily E, member 53 pseudog
 NG_002317 Homo sapiens olfactory receptor, family 7, subfamily E, member 62 pseudog
 NG_002319 Homo sapiens olfactory receptor, family 2, subfamily H, member 5 pseudog
 NG_002320 Homo sapiens NADH dehydrogenase 2 pseudogene 2 (MTND2P2) on chrort
 NG_002321 Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 6 psei
 NG_002322 Homo sapiens olfactory receptor, family 8, subfamily B, member 9 pseudog
 NG_002323 Homo sapiens high mobility group AT-hook 1-like 3 (HMGAI1.3) pseudogene
 NG_002324 Homo sapiens high mobility group AT-hook 1-like 1 (HMGAI1.1) pseudogene
 NG_002325 Homo sapiens COP9 pseudogene (bA345E19.2) on chromosome X
 NG_002326 Homo sapiens dihydroliipoamide S-succinyltransferase pseudogene (E2 com
 NG_002327 Homo sapiens HSA12cenp11 beta-tubulin 4Q (TUBB4Q) pseudogene (LOC
 NG_002328 Homo sapiens polypyrimidine tract binding protein 1 pseudogene (PTBP1)
 NG_002329 Homo sapiens protease, serine, 29 pseudogene (PRSS29P) on chromosom
 NG_002331 Homo sapiens apolipoprotein B mRNA editing enzyme, catalytic polypeptide
 NG_002332 Homo sapiens fascin pseudogene (LOC145989) on chromosome 15
 NG_002333 Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7 psei
 NG_002334 Homo sapiens HSA16q24 beta-tubulin 4Q pseudogene (LOC197331) on chr

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NG_002335 Homo sapiens HSA1q43-44 beta-tubulin 4Q (TUBB4Q) pseudogene (LOC20
 NG_002336 Homo sapiens tubulin, beta polypeptide pseudogene 5 (TUBBP5) on chromo
 NG_002337 Homo sapiens olfactory receptor, family 5, subfamily AK, member 4 pseudog
 NG_002338 Homo sapiens HSA1q42.3 beta-tubulin 4Q pseudogene (LOC255208) on chr
 NG_002339 Homo sapiens HSA18p11 beta-tubulin 4Q pseudogene (LOC280334) on chr
 NG_002340 Homo sapiens ELF2P2 pseudogene (ELF2P2)
 NG_002341 Homo sapiens ELF2P3 pseudogene (AF256221) on chromosome 9
 NG_002342 Homo sapiens zinc finger protein Np97 pseudogene (LOC260337) on chrom
 NG_002343 Homo sapiens HSA12cenq11 beta-tubulin 4Q (TUBB4Q) pseudogene (LOC2
 NG_002348 Homo sapiens sorting nexin 6 pseudogene (LOC126506) on chromosome 15
 NG_002349 Homo sapiens eukaryotic translation elongation factor 1 alpha-like 11 (EEF1
 NG_002350 Homo sapiens eukaryotic translation elongation factor 1 alpha 1 pseudogene
 NG_002351 Homo sapiens sorting nexin 7 pseudogene (LOC203930) on chromosome 1'
 NG_002352 Homo sapiens actin related protein 2/3 complex subunit 1A pseudogene (LO
 NG_002353 Homo sapiens eukaryotic translation elongation factor 1 alpha 1 pseudogene
 NG_002354 Homo sapiens actin related protein 2/3 complex subunit 1A pseudogene (LO
 NG_002355 Homo sapiens actin related protein 2/3 complex subunit 1A pseudogene (LO
 NG_002356 Homo sapiens olfactory receptor, family 7, subfamily E, member 19 pseudog
 NG_002357 Homo sapiens olfactory receptor, family 5, subfamily AH, member 1 pseudog
 NG_002360 Homo sapiens ribosomal protein S2 pseudogene (LOC125208) on chromos
 NG_002361 Homo sapiens CGI-148 protein pseudogene (CGI-148P) on chromosome 16
 NG_002363 Homo sapiens actin related protein 2/3 complex, subunit 3B, 21kDa (ARPC3
 NG_002364 Homo sapiens tumor-associated calcium signal transducer 1 pseudogene (L
 NG_002365 Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) pseudogene
 NG_002366 Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 9, p
 NG_002367 Homo sapiens family with sequence similarity 12, member C pseudogene (F
 NG_002368 Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 3 p
 NG_002369 Homo sapiens Charot-Leyden crystal protein pseudogene 1 (CLCP1) on chr
 NG_002370 Homo sapiens NS1-associated protein 1 pseudogene (LOC149844) on chr
 NG_002371 Homo sapiens dendritic cell protein pseudogene (LOC266683) on chromoso
 NG_002372 Homo sapiens embryonic ectoderm development pseudogene (LOC266694)
 NG_002373 Homo sapiens enhancer of zeste homolog 2 (Drosophila) pseudogene (LOC
 NG_002374 Homo sapiens POM121 membrane glycoprotein-like 4 pseudogene (rat) (PO
 NG_002375 Homo sapiens oligophrenin 1 pseudogene 1 (OPHN1P1) on chromosome 22
 NG_002376 Homo sapiens fatty acid binding protein 5, pseudogene 1 (FABP5P1) on chr
 NG_002378 Homo sapiens ash2 (absent, small, or homeotic)-like (Drosophila) pseudoge
 NG_002379 Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 6 p
 NG_002380 Homo sapiens heat shock 70kDa protein 9B pseudogene (HSPA9BP) on chr
 NG_002381 Homo sapiens ubiquitin protein ligase E3a pseudogene 1 (UBE3AP1) on chr
 NG_002382 Homo sapiens melanoma antigen pseudogene, family A (psMAGEA) on chr
 NG_002383 Homo sapiens keratin 19 pseudogene (LOC160313) on chromosome 12
 NG_002384 Homo sapiens nucleoporin 50 pseudogene (NUP50P) on chromosome 14
 NG_002385 Homo sapiens tropomyosin-like (LOC146253) pseudogene on chromosome
 NG_002387 Homo sapiens proteasome 26S non-ATPase subunit 2 pseudogene (LOC26
 NG_002388 Homo sapiens nucleoporin 50kDa pseudogene (LOC266785) on chromosom
 NG_002389 Homo sapiens nucleoporin 50 kDa pseudogene (LOC266786) on chromosom
 NG_002392 Homo sapiens major histocompatibility complex, class II, DR52 haplotype (D
 NG_002393 Homo sapiens adenylate kinase 2 pseudogene (AK2B) on chromosome 1
 NG_002394 Homo sapiens calcium-binding tyrosine phosphorylation-regulated protein ps
 NG_002395 Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase,
 NG_002397 Homo sapiens major histocompatibility complex, class I, BC (HLA-BC) on chr
 NG_002398 Homo sapiens major histocompatibility complex, class I, GHKAJ (HLA-GHKA
 NG_002399 Homo sapiens transcription elongation factor B (SII), polypeptide 1 pseudog
 NG_002400 Homo sapiens piggyBac transposable element derived 3 pseudogene 1 (PGI
 NG_002401 Homo sapiens piggyBac transposable element derived 3 pseudogene 2 (PGI
 NG_002402 Homo sapiens piggyBac transposable element derived 3 pseudogene 3 (PGI
 NG_002403 Homo sapiens piggyBac transposable element derived 3 pseudogene 4 (PGI

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NG_002404 Homo sapiens ribosomal protein S9 pseudogene 2 (RPS9P2) on chromosom
 NG_002405 Homo sapiens RNA, U12 small nuclear pseudogene (RNU12P) on chromos
 NG_002406 Homo sapiens olfactory receptor, family 5, subfamily G, member 3 pseudog
 NG_002408 Homo sapiens olfactory receptor, family 13, subfamily C, member 7 pseudog
 NG_002409 Homo sapiens olfactory receptor, family 5, subfamily AX, member 1 (OR5AX
 NG_002410 Homo sapiens musashi 1 pseudogene (LOC268276) on chromosome 11
 NG_002411 Homo sapiens suppressor of cytokine signaling 2 pseudogene 1 (SOCS2P1)
 NG_002412 Homo sapiens proteasome 26S non-ATPase subunit 7 pseudogene (LOC281
 NG_002415 Homo sapiens RNA binding motif protein 8B pseudogene (RBM8B) on chrom
 NG_002416 Homo sapiens mitochondrial ribosomal protein L11 pseudogene (MRPL11P2)
 NG_002417 Homo sapiens proteasome 26S non-ATPase subunit 10 pseudogene (LOC2
 NG_002418 Homo sapiens similar to Homeobox protein Meis3 (Meis1-related protein 2) (
 NG_002419 Homo sapiens eukaryotic translation initiation factor 1A pseudogene 1 (EIF1;
 NG_002423 Homo sapiens CD8 antigen, beta polypeptide 2, pseudogene (p37) (CD8B2)
 NG_002425 Homo sapiens RNA, 7SL, cytoplasmic, pseudogene 1 (RN7SLP1)
 NG_002426 Homo sapiens RNA, 7SL, cytoplasmic, pseudogene 2 (RN7SLP2) on chrom
 NG_002427 Homo sapiens RNA, 7SL, cytoplasmic, pseudogene 3 (RN7SLP3)
 NG_002428 Homo sapiens RNA, 7SL, cytoplasmic, pseudogene 4 (RN7SLP4) on chrom
 NG_002429 Homo sapiens RNA, 7SL, cytoplasmic, pseudogene 5 (RN7SLP5) on chrom
 NG_002430 Homo sapiens mitochondrial ribosomal protein L48 pseudogene 1 (MRPL48
 NG_002431 Homo sapiens nuclear autoantigenic sperm protein pseudogene 1 (NASPP1;
 NG_002432 Homo sapiens major histocompatibility complex, class II, DR51 haplotype (D
 NG_002433 Homo sapiens major histocompatibility complex, class II, DR53 haplotype (D
 NG_002434 Homo sapiens pre-B-cell leukemia transcription factor pseudogene 1 (PBXP
 NG_002437 Homo sapiens cytochrome b-5 pseudogene 3 (CYB5P3) on chromosome 14
 NG_002438 Homo sapiens LEFTY family pseudogene (LEFTY3) on chromosome 1
 NG_002440 Homo sapiens TGFB-induced factor (TALE family homeobox) pseudogene (l
 NG_002448 Homo sapiens mitogen-activated protein kinase 6 pseudogene 2 (MAPK6PS
 NG_002449 Homo sapiens mitogen-activated protein kinase 6 pseudogene 6 (MAPK6PS
 NG_002450 Homo sapiens cytokine receptor-like factor 3 pseudogene (LOC285706) on c
 NG_002451 Homo sapiens mitogen-activated protein kinase 6 pseudogene 4 (MAPK6PS
 NG_002452 Homo sapiens mitogen-activated protein kinase 6 pseudogene 5 (MAPK6PS
 NG_002453 Homo sapiens mitogen-activated protein kinase 6 pseudogene 3 (MAPK6PS
 NG_002454 Homo sapiens mitogen-activated protein kinase 6 pseudogene 1 (MAPK6PS
 NG_002456 Homo sapiens ribosomal protein L32-like 2 (RPL32L2) pseudogene on chro
 NG_002457 Homo sapiens protein tyrosine phosphatase, non-receptor type substrate 1-l
 NG_002458 Homo sapiens actin, beta-like 1 (ACTBL1) pseudogene on chromosome 22
 NG_002459 Homo sapiens vomeronasal 1 receptor 9 pseudogene (VN1R9P) on chromos
 NG_002460 Homo sapiens high-mobility group nucleosomal binding domain 2 pseudog
 NG_002461 Homo sapiens ataxin 2 related protein pseudogene (LOC317727) on chromo
 NG_002462 Homo sapiens basic leucine zipper nuclear factor 2 pseudogene (BLZF2P) o
 NG_002463 Homo sapiens proteasome 26S non-ATPase subunit 12 pseudogene (LOC3
 NG_002464 Homo sapiens steroidogenic acute regulator pseudogene 1 (STARP1) on chi
 NG_002465 Homo sapiens chromosome 14 open reading frame 55 (C14orf55) pseudoge
 NG_002467 Homo sapiens a disintegrin and metalloproteinase domain 21 pseudogene (/
 NG_002468 Homo sapiens ribosomal protein L9 pseudogene (LOC254948) on chromoso
 NG_002469 Homo sapiens bromodomain containing 7 pseudogene (BRD7P) on chromos
 NG_002470 Homo sapiens basic transcription factor 3, pseudogene 2 (BTF3P2) on chro
 NG_002471 Homo sapiens ribosomal protein L9 pseudogene (LOC317771) on chromoso
 NG_002472 Homo sapiens CDC10 cell division cycle 10 homolog (S. cerevisiae) pseudog
 NG_002473 Homo sapiens cysteine and histidine-rich domain (CHORD)-containing 2 pse
 NG_002474 Homo sapiens CNDBP1 interactor pseudogene (CBPINP) on chromosom
 NG_002475 Homo sapiens CDC28 protein kinase regulatory subunit 1B pseudogene (CK
 NG_002476 Homo sapiens coilin pseudogene (COILP) on chromosome 14
 NG_002477 Homo sapiens cytochrome c oxidase subunit Va pseudogene 2 (COX5AP2) ;
 NG_002478 Homo sapiens ras homolog gene family, member Q pseudogene (RHQQP) c
 NG_002479 Homo sapiens cytochrome c oxidase subunit VIIa polypeptide 3 pseudogene

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NG_002480 Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 18 pseudogene 1 (l
 NG_002481 Homo sapiens G protein-coupled receptor 57 (GPR57) pseudogene on chrom
 NG_002482 Homo sapiens ribosomal protein L21 pseudogene 12 (RPL21P12) on chrom
 NG_002483 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) pseudogene 4 (PPI
 NG_002484 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) pseudogene 5 (PPI
 NG_002485 Homo sapiens Friedrich ataxia pseudogene (FRDAP) on chromosome 14
 NG_002486 Homo sapiens laminin receptor 1 pseudogene 4 (LAMR1P4) on chromosom
 NG_002487 Homo sapiens ribosomal protein L21 pseudogene 7 (RPL21P7) on chromos
 NG_002488 Homo sapiens ribosomal protein L21 pseudogene 11 (RPL21P11) on chromi
 NG_002489 Homo sapiens laminin receptor 1 pseudogene 3 (LAMR1P3) on chromosom
 NG_002490 Homo sapiens ubiquitin-conjugating enzyme E2L 7 pseudogene (UBE2L7) o
 NG_002491 Homo sapiens ribosomal protein L21 pseudogene 9 (RPL21P9) on chromos
 NG_002492 Homo sapiens Rho GTPase activating protein 16 pseudogene (ARHGAP16F
 NG_002493 Homo sapiens DnaJ (Hsp40) homolog, subfamily C, member 8 pseudogene
 NG_002494 Homo sapiens eukaryotic translation elongation factor 1 alpha 1 pseudogene
 NG_002495 Homo sapiens endosulfine alpha pseudogene 2 (ENSAP2) on chromosome
 NG_002496 Homo sapiens eukaryotic translation initiation factor 2, subunit 2 beta, pseud
 NG_002497 Homo sapiens ATPase, H⁺ transporting, lysosomal 13kDa, V1 subunit G pse
 NG_002498 Homo sapiens eukaryotic translation initiation factor 3, subunit 6 interacting f
 NG_002499 Homo sapiens eukaryotic translation initiation factor 4B pseudogene (EIF4B)
 NG_002500 Homo sapiens ATP synthase, H⁺ transporting, mitochondrial F0 complex, su
 NG_002501 Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase acti
 NG_002502 Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase acti
 NG_002503 Homo sapiens uracil-DNA glycosylase pseudogene 3 (UNG3P3) on chromosc
 NG_002504 Homo sapiens zinc finger protein 405 pseudogene (ZNF405P) on chromosom
 NG_002505 Homo sapiens asparaginyl-tRNA synthetase pseudogene (NARSP) on chrom
 NG_002506 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) pseudogene 6 (PPI
 NG_002507 Homo sapiens laminin receptor 1 pseudogene 5 (LAMR1P5) on chromosom
 NG_002508 Homo sapiens high-mobility group box 1 pseudogene (HMGB1P) on chromo
 NG_002509 Homo sapiens high-mobility group nucleosome binding domain 1 pseudogen
 NG_002510 Homo sapiens high-mobility group nucleosome binding domain 2 pseudogen
 NG_002511 Homo sapiens heterogeneous nuclear ribonucleoprotein C pseudogene (HNI
 NG_002512 Homo sapiens heterogeneous nuclear ribonucleoprotein U pseudogene (HNI
 NG_002513 Homo sapiens ribosomal protein L10a pseudogene 1 (RPL10AP1) on chrom
 NG_002514 Homo sapiens ATP synthase, H⁺ transporting, mitochondrial F0 complex, su
 NG_002515 Homo sapiens ATP synthase, H⁺ transporting, mitochondrial F0 complex, su
 NG_002516 Homo sapiens BCL2/adenovirus E1B 19kDa interacting protein 3 pseudogen
 NG_002517 Homo sapiens glycine C-acetyltransferase pseudogene (GCATP) on chromo
 NG_002518 Homo sapiens ribosomal protein L12 pseudogene 5 (RPL12P5) on chromos
 NG_002519 Homo sapiens ribosomal protein L21 pseudogene 10 (RPL21P10) on chromi
 NG_002520 Homo sapiens ribosomal protein L21 pseudogene 13 (RPL21P13) on chromi
 NG_002521 Homo sapiens ribosomal protein L21 pseudogene 8 (RPL21P8) on chromos
 NG_002522 Homo sapiens ribosomal protein S2 pseudogene 4 (RPS2P4) on chromosom
 NG_002523 Homo sapiens ribosomal protein S29 pseudogene 1 (RPS29P1) on chromos
 NG_002524 Homo sapiens ubiquitin A-52 residue ribosomal protein fusion product 1 pse
 NG_002525 Homo sapiens ribosomal protein L24 pseudogene 3 (RPL24P3) on chromos
 NG_002526 Homo sapiens ribosomal protein, large, P1 pseudogene 1 (RPLP1P1) on chr
 NG_002527 Homo sapiens ribosomal protein L26 pseudogene 4 (RPL26P4) on chromos
 NG_002528 Homo sapiens ubiquitin-conjugating enzyme E2C pseudogene 1 (UBE2CP1)
 NG_002529 Homo sapiens tubulin, beta polypeptide pseudogene 3 (TUBBP3) on chromo
 NG_002530 Homo sapiens serine/threonine kinase 16 pseudogene (STK16P) on chromo
 NG_002531 Homo sapiens abl-interactor 1 pseudogene (ABI1P) on chromosome 14
 NG_002532 Homo sapiens spermidine synthase pseudogene 2 (SRMP2) on chromosom
 NG_002533 Homo sapiens small nuclear ribonucleoprotein polypeptide G pseudogene (S
 NG_002534 Homo sapiens solute carrier family 20 (phosphate transporter), member 1 ps
 NG_002535 Homo sapiens ribosomal protein L12 pseudogene 7 (RPL12P7) on chromosc
 NG_002537 Homo sapiens SET pseudogene 2 (SETP2) on chromosome 14

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NG_002538 Homo sapiens ribosomal protein L13a pseudogene 2 (RPL13AP2) on chrom
 NG_002539 Homo sapiens ribosomal protein L15 pseudogene 2 (RPL15P2) on chromos
 NG_002540 Homo sapiens ribosomal protein L17 pseudogene 3 (RPL17P3) on chromos
 NG_002541 Homo sapiens ribosomal protein L17 pseudogene 4 (RPL17P4) on chromos
 NG_002542 Homo sapiens ribosomal protein L18a pseudogene 1 (RPL18AP1) on chrom
 NG_002543 Homo sapiens ribosomal protein L18 pseudogene 1 (RPL18P1) on chromos
 NG_002544 Homo sapiens ribosomal protein L22 pseudogene 2 (RPL22P2) on chromos
 NG_002545 Homo sapiens ribosomal protein L23a pseudogene 10 (RPL23AP10) on chr
 NG_002546 Homo sapiens ribosomal protein L23a pseudogene 11 (RPL23AP11) on chr
 NG_002547 Homo sapiens eukaryotic translation initiation factor 4E binding protein 1 pse
 NG_002548 Homo sapiens heat shock factor binding protein 1 pseudogene 1 (HSBP1P1)
 NG_002549 Homo sapiens ribosomal protein L26 pseudogene 2 (RPL26P2) on chromos
 NG_002550 Homo sapiens ribosomal protein L26 pseudogene 3 (RPL26P3) on chromos
 NG_002551 Homo sapiens ribosomal protein L27 pseudogene 1 (RPL27P1) on chromos
 NG_002552 Homo sapiens heat shock 10kDa protein 1 (chaperonin 10) pseudogene 2 (H
 NG_002553 Homo sapiens ribosomal protein L36a pseudogene 2 (RPL36AP2) on chrom
 NG_002554 Homo sapiens NEK2 pseudogene (NEK2P) on chromosome 14
 NG_002555 Homo sapiens ribosomal protein L36a pseudogene 3 (RPL36AP3) on chrom
 NG_002556 Homo sapiens ribosomal protein L36a pseudogene 4 (RPL36AP4) on chrom
 NG_002557 Homo sapiens molybdenum cofactor synthesis 3 pseudogene (MOC53P) on chrom
 NG_002558 Homo sapiens ribosomal protein L39 pseudogene 2 (RPL39P2) on chromos
 NG_002559 Homo sapiens ribosomal protein L3 pseudogene 4 (RPL3P4) on chromosome
 NG_002560 Homo sapiens ribosomal protein L41 pseudogene 4 (RPL41P4) on chromos
 NG_002561 Homo sapiens ribosomal protein L7a pseudogene 5 (RPL7AP5) on chromos
 NG_002562 Homo sapiens ribosomal protein L7a pseudogene 6 (RPL7AP6) on chromos
 NG_002563 Homo sapiens ribosomal protein L9 pseudogene 6 (RPL9P6) on chromosome
 NG_002564 Homo sapiens ribosomal protein S12 pseudogene 1 (RPS12P1) on chromos
 NG_002565 Homo sapiens pituitary tumor-transforming 4 pseudogene (PTTG4P) on chr
 NG_002566 Homo sapiens ribosomal protein S15a pseudogene 2 (RPS15AP2) on chrom
 NG_002567 Homo sapiens ribosomal protein S15a pseudogene 3 (RPS15AP3) on chrom
 NG_002568 Homo sapiens olfactory receptor, family 11, subfamily H, member 8 pseudog
 NG_002570 Homo sapiens ribosomal protein S18 pseudogene 2 (RPS18P2) on chromos
 NG_002571 Homo sapiens ribosomal protein S24 pseudogene 2 (RPS24P2) on chromos
 NG_002572 Homo sapiens ribosomal protein S24 pseudogene 3 (RPS24P3) on chromos
 NG_002573 Homo sapiens ribosomal protein S27a pseudogene 4 (RPS27AP4) on chrom
 NG_002574 Homo sapiens ribosomal protein S8 pseudogene 1 (RPS8P1) on chromosome
 NG_002575 Homo sapiens ribosomal protein S2 pseudogene 2 (RPS2P2) on chromosome
 NG_002576 Homo sapiens ribosomal protein S2 pseudogene 3 (RPS2P3) on chromosome
 NG_002577 Homo sapiens SSX1 pseudogene (psiSSX1) on chromosome X
 NG_002578 Homo sapiens SSX4 pseudogene (psiSSX4) on chromosome X
 NG_002579 Homo sapiens SSX5 pseudogene (psiSSX5) on chromosome X
 NG_002580 Homo sapiens SSX3 pseudogene (psiSSX3) on chromosome X
 NG_002581 Homo sapiens mago-nashi homolog, proliferation-associated pseudogene (C
 NG_002582 Homo sapiens SSX2 pseudogene (psiSSX2) on chromosome X
 NG_002583 Homo sapiens psiSSX8 pseudogene (psiSSX8) on chromosome X
 NG_002584 Homo sapiens SSX9 pseudogene (psiSSX9) on chromosome X
 NG_002585 Homo sapiens SSX6 pseudogene (psiSSX6) on chromosome X
 NG_002586 Homo sapiens SSX10 pseudogene (psiSSX10) on chromosome 6
 NG_002587 Homo sapiens SSX7 pseudogene (psiSSX7) on chromosome X
 NG_002588 Homo sapiens mortality factor 4 like pseudogene 1 (MORF4LP1) on chromos
 NG_002589 Homo sapiens mortality factor 4 like pseudogene 2 (MORF4LP2) on chromos
 NG_002590 Homo sapiens mortality factor 4 like pseudogene 3 (MORF4LP3) on chromos
 NG_002591 Homo sapiens mortality factor 4 like pseudogene 4 (MORF4LP4) on chromos
 NG_002592 Homo sapiens MAD2 mitotic arrest deficient-like 1 (yeast) pseudogene (MAC
 NG_002593 Homo sapiens cytochrome c oxidase pseudogene 2 (MTCO1P2) on chromos
 NG_002594 Homo sapiens nuclear receptor coactivator 4 pseudogene (NCOA4P) on chr
 NG_002595 Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8, ps

NG_002596 Homo sapiens nicotinamide nucleotide adenyltransferase pseudogene (NM
 NG_002597 Homo sapiens nucleophosmin 1 (nucleolar phosphoprotein B23, numatrin) p
 NG_002598 Homo sapiens 5',3'-nucleotidase, cytosolic pseudogene 1 (NT5CP1) on chro
 NG_002599 Homo sapiens 5',3'-nucleotidase, cytosolic pseudogene 2 (NT5CP2) on chro
 NG_002601 Homo sapiens UDP glycosyltransferase 1 family, polypeptide A cluster (UGT
 NG_002603 Homo sapiens nuclear transport factor 2 pseudogene 2 (NUTF2P2) on chro
 NG_002604 Homo sapiens prostatic binding protein pseudogene 1 (PBPP1) on chromos
 NG_002605 Homo sapiens PCQAP pseudogene (PCQAPP) on chromosome 14
 NG_002606 Homo sapiens PDZ and LIM domain 1 pseudogene (PDLIM1P) on chromos
 NG_002607 Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 3 pse
 NG_002608 Homo sapiens prothymosin, alpha pseudogene 7 (PTMAP7) on chromosom
 NG_002609 Homo sapiens RAN binding protein 20 pseudogene (RANBP20P) on chromo
 NG_002610 Homo sapiens replication protein A2 pseudogene (RPA2P) on chromosome
 NG_002611 Homo sapiens interleukin 6 signal transducer (gp130, oncostatin M receptor)
 NG_002612 Homo sapiens family with sequence similarity 16, member A, Y-linked (FAM1
 NG_002613 Homo sapiens family with sequence similarity 16, member B (FAM16B) pseu
 NG_002616 Homo sapiens histone 2, H3b (HIST2H3B) pseudogene on chromosome 1
 NG_002617 Homo sapiens histone 2, H3a (HIST2H3A) pseudogene on chromosome 1
 NG_002618 Homo sapiens histone 3, H2ba (HIST3H2BA) pseudogene on chromosome 1
 NG_002619 Homo sapiens histone 2, H2bc (HIST2H2BC) pseudogene on chromosome 1
 NG_002620 Homo sapiens histone 2, H2bd (HIST2H2BD) pseudogene on chromosome 1
 NG_002621 Homo sapiens histone 2, H2ba (HIST2H2BA) pseudogene on chromosome 1
 NG_002622 Homo sapiens ELYS transcription factor-like protein TMB52 pseudogene (L
 NG_002623 Homo sapiens ADP-ribosyltransferase (NAD⁺: poly (ADP-ribose) polymerase
 NG_002624 Homo sapiens adaptin, beta 1-like 2 (ADTB1L2) pseudogene on chromosom
 NG_002625 Homo sapiens adaptin, beta 1-like 1 (ADTB1L1) pseudogene on chromosom
 NG_002626 Homo sapiens proteasome (prosome, macropain) activator subunit 2 pseud
 NG_002627 Homo sapiens proteasome 26S non-ATPase subunit 10 pseudogene (LOC3:
 NG_002628 Homo sapiens proteasome activator subunit 2 pseudogene (LOC338095) on
 NG_002629 Homo sapiens proteasome activator subunit 2 pseudogene (LOC338096) on
 NG_002630 Homo sapiens proteasome activator subunit 2 pseudogene (LOC338097) on
 NG_002631 Homo sapiens proteasome activator subunit 2 pseudogene (LOC338098) on
 NG_002632 Homo sapiens proteasome activator subunit 2 pseudogene (LOC338099) on
 NG_002633 Homo sapiens cytochrome c oxidase subunit Vb-like 7 (COX5BL7) pseudoge
 NG_002634 Homo sapiens serum amyloid A3 pseudogene (SAA3P) on chromosome 11
 NG_002636 Homo sapiens YME1-like 2 (S. cerevisiae) (YME1L2) pseudogene on chrom
 NG_002637 Homo sapiens argininosuccinate lyase-like (ASLL) pseudogene on chromos
 NG_002638 Homo sapiens colony stimulating factor 2 receptor, beta, 2 (CSF2RB2) pseu
 NG_002639 Homo sapiens inorganic pyrophosphatase pseudogene (LOC151842) on chr
 NG_002640 Homo sapiens mannan-binding lectin serine protease 1 pseudogene 1 (MAS
 NG_002641 Homo sapiens inorganic pyrophosphatase pseudogene (LOC285591) on chr
 NG_002642 Homo sapiens SUL1D pseudogene (SUL1D1P) on chromosome 4
 NG_002645 Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin),
 NG_002646 Homo sapiens high-mobility group (nonhistone chromosomal) protein 17-like
 NG_002647 Homo sapiens vomeronasal 1 receptor 8 pseudogene (VN1R8P) on chromo:
 NG_002648 Homo sapiens mitochondrial ribosomal protein S16 pseudogene (MRPS16P
 NG_002649 Homo sapiens mitochondrial ribosomal protein S16 pseudogene (MRPS16P
 NG_002650 Homo sapiens mitochondrial ribosomal protein S24 pseudogene (MRPS24P
 NG_002651 Homo sapiens mitochondrial ribosomal protein S25 pseudogene (MRPS25P
 NG_002652 Homo sapiens histone 2, H2bb (HIST2H2BB) pseudogene on chromosome 5
 NG_002653 Homo sapiens taste receptor, type 2, member 62 pseudogene (TAS2R62P)
 NG_002655 Homo sapiens ADP-ribosyltransferase (NAD⁺: poly (ADP-ribose) polymerase
 NG_002656 Homo sapiens ADP-ribosyltransferase (NAD⁺: poly (ADP-ribose) polymerase
 NG_002657 Homo sapiens DEAH (Asp-Glu-Ala-His) box polypeptide 9 pseudogene (DH)
 NG_002658 Homo sapiens eukaryotic translation initiation factor 4E-like 2 (EIF4E12) pse
 NG_002659 Homo sapiens zinc finger pseudogene (BA393J16.4) on chromosome 10
 NG_002660 Homo sapiens KRAB box zinc finger protein pseudogene (BA775A3.1) on ch

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NG_002661 Homo sapiens hypothetical pseudogene bA291L22.4 (bA291L22.4) on chrom
 NG_002662 Homo sapiens mitochondrial ribosomal protein S5 pseudogene (MRPS5P3)
 NG_002663 Homo sapiens mitochondrial ribosomal protein S21 pseudogene (MRPS21P)
 NG_002665 Homo sapiens mitochondrial ribosomal protein L42 pseudogene (MRPL42P1)
 NG_002666 Homo sapiens mitochondrial ribosomal protein S18C pseudogene (MRPS18C)
 NG_002667 Homo sapiens keratin associated protein 13 pseudogene 1 (KRTAP13P1) or
 NG_002668 Homo sapiens keratin associated protein 13 pseudogene 2 (KRTAP13P2) or
 NG_002669 Homo sapiens keratin associated protein 8 pseudogene 1 (KRTAP8P1) on ch
 NG_002670 Homo sapiens keratin associated protein 8 pseudogene 2 (KRTAP8P2) on ch
 NG_002671 Homo sapiens keratin associated protein 19 pseudogene 3 (KRTAP19P3) or
 NG_002672 Homo sapiens keratin associated protein 19 pseudogene 4 (KRTAP19P4) or
 NG_002673 Homo sapiens keratin associated protein 21 pseudogene 1 (KRTAP21P1) or
 NG_002674 Homo sapiens taste receptor, type 2, member 64 pseudogene (TAS2R64P)
 NG_002675 Homo sapiens taste receptor, type 2, member 63 pseudogene (TAS2R63P)
 NG_002676 Homo sapiens taste receptor, type 2, member 65 pseudogene (TAS2R65P)
 NG_002679 Homo sapiens potassium large conductance calcium-activated channel, subf
 NG_002680 Homo sapiens survival motor neuron pseudogene (SMN) on chromosome 5
 NG_002681 Homo sapiens leukocyte immunoglobulin-like receptor, subfamily A (without
 NG_002682 Homo sapiens MAPRE1P pseudogene (MAPRE1P) on chromosome 8
 NG_002683 Homo sapiens steroid-5-beta-reductase, beta polypeptide pseudogene 1 (SF
 NG_002684 Homo sapiens mitochondrial translational initiation factor 2 pseudogene 1 (M
 NG_002685 Homo sapiens histone 1, H2a, pseudogene 2 (HIST1H2APS2) on chromoso
 NG_002687 Homo sapiens suppressor of cytokine signaling 2 pseudogene 2 (SOCS2P2)
 NG_002688 Homo sapiens keratin associated protein 19 pseudogene 1 (KRTAP19P1) or
 NG_002689 Homo sapiens keratin associated protein 19 pseudogene 2 (KRTAP19P2) or
 NG_002690 Homo sapiens Prader-Willi/Angelman syndrome region (PWSAS) on chrom
 NG_002691 Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, ep
 NG_002692 Homo sapiens protein phosphatase 1A pseudogene (LOC137012) on chrom
 NG_002693 Homo sapiens phosphoribosyl pyrophosphate amidotransferase pseudogene
 NG_002694 Homo sapiens mesoderm specific transcript homolog (mouse) pseudogene (H
 NG_002696 Homo sapiens eukaryotic translation initiation factor 2, subunit 2 beta, pseud
 NG_002697 Homo sapiens ALEX2 pseudogene (LOC347674) on chromosome 7
 NG_002698 Homo sapiens eukaryotic translation initiation factor 2 beta-like pseudogene
 NG_002699 Homo sapiens G protein gamma 5-like subunit (GN5ps) pseudogene on ch
 NG_002700 Homo sapiens endothelin converting enzyme-like 1, pseudogene 1 (ECE1P
 NG_002701 Homo sapiens ECEL2 pseudogene 2 (ECEL2) on chromosome 2
 NG_002702 Homo sapiens calgizzarin-like (LOC347701) pseudogene on chromosome 7
 NG_002703 Homo sapiens mitochondrial ribosomal protein S38 pseudogene (MRPS38P)
 NG_002705 Homo sapiens mitochondrial ribosomal protein S36 pseudogene (MRPS36P)
 NG_002707 Homo sapiens mitochondrial ribosomal protein S36 pseudogene (MRPS36P)
 NG_002709 Homo sapiens mitochondrial ribosomal protein S36 pseudogene (MRPS36P)
 NG_002711 Homo sapiens mitochondrial ribosomal protein S36 pseudogene (MRPS36P)
 NG_002713 Homo sapiens mitochondrial ribosomal protein S36 pseudogene (MRPS36P)
 NG_002716 Homo sapiens calmodulin 2 pseudogene 1 (CALM2P1) on chromosome 17
 NG_002717 Homo sapiens nitric oxide synthase 2A pseudogene (LOC284193) on chrom
 NG_002718 Homo sapiens galectin-9 pseudogene (LOC284194) on chromosome 17
 NG_002719 Homo sapiens TL132 pseudogene (LOC347716) on chromosome 17
 NG_002720 Homo sapiens signal recognition particle 68kD pseudogene (LOC347717) or
 NG_002721 Homo sapiens karyopherin (importin) beta 2 pseudogene (LOC347719) on ch
 NG_002723 Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase,
 NG_002724 Homo sapiens actin, beta pseudogene 7 (ACTBP7) on chromosome 15
 NG_002725 Homo sapiens eukaryotic translation initiation factor 5A pseudogene 3 (EIF5
 NG_002726 Homo sapiens SRY (sex determining region Y)-box 5 pseudogene (SOX5P)
 NG_002727 Homo sapiens MHC class I polypeptide-related sequence C (MICC) pseudog
 NG_002728 Homo sapiens heterogeneous nuclear ribonucleoprotein A1 pseudogene (hn
 NG_002729 Homo sapiens HLA-75 pseudogene (HLA-75) on chromosome 6
 NG_002731 Homo sapiens HLA-90 pseudogene (HLA-90) on chromosome 6

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NG_002733 Homo sapiens MHC class I polypeptide-related sequence G pseudogene (Ml
 NG_002735 Homo sapiens 3.8-1.5 pseudogene (3.8-1.5) on chromosome 6
 NG_002736 Homo sapiens HLA complex P5 pseudogene 12 (HCP5P12) on chromosome
 NG_002737 Homo sapiens HLA complex group 2 pseudogene 8 (HCG2P8) on chromoso
 NG_002738 Homo sapiens ribosomal protein L7a pseudogene 7 (RPL7AP7) on chromos
 NG_002739 Homo sapiens HLA complex group 4 pseudogene 9 (HCG4P9) on chromoso
 NG_002740 Homo sapiens HLA complex P5 pseudogene 13 (HCP5P13) on chromosome
 NG_002741 Homo sapiens HLA complex group 4 pseudogene 10 (HCG4P10) on chromo
 NG_002742 Homo sapiens HLA complex P5 pseudogene 14 (HCP5P14) on chromosome
 NG_002743 Homo sapiens HLA complex group 9 pseudogene 5 (HCG9P5) on chromoso
 NG_002744 Homo sapiens HLA complex group 4 pseudogene 11 (HCG4P11) on chromo
 NG_002745 Homo sapiens HLA complex P5 pseudogene 15 (HCP5P15) on chromosome
 NG_002746 Homo sapiens eukaryotic translation initiation factor 5A pseudogene 2 (EIF5
 NG_002747 Homo sapiens hydroxysteroid (17-beta) dehydrogenase pseudogene 1 (HSD
 NG_002748 Homo sapiens interleukin 6 signal transducer (gp130, oncostatin M receptor)
 NG_002749 Homo sapiens RAN, member RAS oncogene family pseudogene 1 (RANP1)
 NG_002750 Homo sapiens SMT3 suppressor of mi1 two 3 homolog 2 (yeast) pseudogene
 NG_002752 Homo sapiens NOD24 pseudogene (NOD24) on chromosome X
 NG_002753 Homo sapiens NOD13 pseudogene (NOD13) on chromosome X
 NG_002754 Homo sapiens NOD25 pseudogene (NOD25) on chromosome 12
 NG_002761 Homo sapiens complement component 4 binding protein, alpha-like 2 (C4BP
 NG_002762 Homo sapiens crystallin, gamma E pseudogene 1 (CRYGEP1) on chromoso
 NG_002763 Homo sapiens glutamate dehydrogenase pseudogene 3 (GLUDP3) on chrom
 NG_002764 Homo sapiens interleukin 9 receptor pseudogene 4 (IL9RP4) on chromosom
 NG_002765 Homo sapiens prothymosin, alpha pseudogene 2 (gene sequence 32) (PTM/
 NG_002766 Homo sapiens cytochrome P450, subfamily 51 pseudogene 2 (CYP51P2) on
 NG_002767 Homo sapiens glycoprotein, alpha-galactosyltransferase 1 pseudogene (GG
 NG_002768 Homo sapiens beta-lactoglobulin pseudogene (LOC138159) on chromosome
 NG_002769 Homo sapiens heat shock 70kDa protein 8 pseudogene (LOC158714) on chi
 NG_002770 Homo sapiens pseudogene of IGF-II mRNA-binding protein 3 (LOC346296) c
 NG_002771 Homo sapiens major histocompatibility complex, class I, L (HLA-L) pseudoge
 NG_002772 Homo sapiens CAP, adenylate cyclase-associated protein, 2 (yeast) pseudog
 NG_002773 Homo sapiens coactosin-like 1 (Dictyostellum) pseudogene 2 (COTL1P2) on
 NG_002775 Homo sapiens keratin pseudogene (LOC147228) on chromosome 17
 NG_002776 Homo sapiens keratin pseudogene (LOC284198) on chromosome 17
 NG_002777 Homo sapiens keratin pseudogene (LOC339186) on chromosome 17
 NG_002778 Homo sapiens keratin pseudogene (LOC339241) on chromosome 17
 NG_002779 Homo sapiens keratin pseudogene (LOC339244) on chromosome 17
 NG_002780 Homo sapiens keratin pseudogene (LOC339258) on chromosome 17
 NG_002781 Homo sapiens keratin pseudogene (LOC353194) on chromosome 17
 NG_002782 Homo sapiens keratin pseudogene (LOC353196) on chromosome 17
 NG_002785 Homo sapiens interleukin 6 receptor pseudogene (LOC157916) on chromoso
 NG_002786 Homo sapiens makorin, ring finger protein, pseudogene 2 (MKRNP2) on chr
 NG_002787 Homo sapiens chromosome 20 open reading frame 189 (C20orf189) pseudo
 NG_002788 Homo sapiens bitter taste receptor pseudogene 8 (PS8) on chromosome 12
 NG_002790 Homo sapiens pseudogene of origin recognition complex, subunit 1-like (LOC
 NG_002791 Homo sapiens selenoprotein W, 1 pseudogene (SEPW1P) on chromosome
 NG_002792 Homo sapiens PC4 and SFRS1 interacting protein 1 pseudogene (PSIP1P) c
 NG_002793 Homo sapiens pelota/integrin, alpha 1 region (PELO/ITGA1@) on chromosom
 NG_002795 Homo sapiens polycystic kidney disease 1 (autosomal dominant) pseudogen
 NG_002796 Homo sapiens polycystic kidney disease 1 (autosomal dominant) pseudogen
 NG_002797 Homo sapiens polycystic kidney disease 1 (autosomal dominant) pseudogen
 NG_002798 Homo sapiens polycystic kidney disease 1 (autosomal dominant) pseudogen
 NG_002799 Homo sapiens polycystic kidney disease 1 (autosomal dominant) pseudogen
 NG_002800 Homo sapiens polycystic kidney disease 1 (autosomal dominant) pseudogen
 NG_002801 Homo sapiens 18S ribosomal RNA pseudogene (LOC359724) on chromosom
 NG_002802 Homo sapiens ADP-ribosyltransferase (NAD+; poly (ADP-ribose) polymerase

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NG_002803 Homo sapiens apical protein-like (Xenopus laevis) pseudogene (APXLP) on c
 NG_002804 Homo sapiens arylsulfatase F pseudogene (ARSFP)
 NG_002805 Homo sapiens calcium/calmodulin-dependent serine protein kinase (MAGUK
 NG_002806 Homo sapiens adlcan pseudogene (ADLICANP) on chromosome Y
 NG_002807 Homo sapiens cofactor required for Sp1 transcriptional activation, subunit 2,
 NG_002808 Homo sapiens C-terminal binding protein 2 pseudogene (LOC352905) on ch
 NG_002809 Homo sapiens eukaryotic translation initiation factor 4A, isoform 1 pseudoge
 NG_002810 Homo sapiens hypothetical protein FLJ10842 pseudogene (LOC359793) on
 NG_002811 Homo sapiens glycogenin 2 pseudogene (GYG2P) on chromosome Y
 NG_002812 Homo sapiens lung cancer candidate FUS1 pseudogene (LOC359794) on c
 NG_002813 Homo sapiens neurofilament, light polypeptide 68kDa pseudogene (LOC359
 NG_002814 Homo sapiens G protein-coupled receptor 143 pseudogene (GPR143P) on c
 NG_002815 Homo sapiens 60S ribosomal protein L26 pseudogene (LOC347593) on chr
 NG_002816 Homo sapiens splicing factor proline/glutamine rich (polypyrimidine tract binc
 NG_002817 Homo sapiens solute carrier family 25 (mitochondrial carrier; ornithine transp
 NG_002818 Homo sapiens Smcy homolog, Y chromosome (mouse) pseudogene (SMCYI
 NG_002819 Homo sapiens transducin (beta)-like 1Y-linked pseudogene (TBL1YP) on chr
 NG_002821 Homo sapiens ubiquitin-conjugating enzyme E2 variant 1 pseudogene (LOC:
 NG_002822 Homo sapiens ubiquitin specific protease 12 pseudogene 1 (USP12P1) on d
 NG_002823 Homo sapiens ubiquitin specific protease 12 pseudogene 2 (USP12P2) on d
 NG_002824 Homo sapiens ubiquitin specific protease 12 pseudogene 3 (USP12P3) on d
 NG_002825 Homo sapiens voltage-dependent anion channel 1 pseudogene (LOC359800
 NG_002826 Homo sapiens Ras-homolog enriched in brain pseudogene 1 (RHEBP1) on c
 NG_002827 Homo sapiens mitochondrial ribosomal protein 63 pseudogene 1 (MRP63P1;
 NG_002828 Homo sapiens mitochondrial ribosomal protein S5 pseudogene (LOC133332
 NG_002829 Homo sapiens mitochondrial ribosomal protein 63 pseudogene 6 (MRP63P6;
 NG_002830 Homo sapiens mitochondrial ribosomal protein S33 pseudogene 1 (MRP533
 NG_002831 Homo sapiens mitochondrial ribosomal protein L36 pseudogene 1 (MRPL361
 NG_002832 Homo sapiens mitochondrial ribosomal protein S35 pseudogene 1 (MRP535
 NG_002833 Homo sapiens mitochondrial ribosomal protein S7 pseudogene 2 (MRPS7P2
 NG_002834 Homo sapiens mitochondrial ribosomal protein L51 pseudogene 2 (MRPL511
 NG_002835 Homo sapiens mitochondrial ribosomal protein S18C pseudogene 6 (MRPS1
 NG_002836 Homo sapiens mitochondrial ribosomal protein L49 pseudogene 2 (MRPL491
 NG_002837 Homo sapiens mitochondrial ribosomal protein 63 pseudogene 10 (MRP63P
 NG_002838 Homo sapiens mitochondrial ribosomal protein L2 pseudogene 1 (MRPL2P1;
 NG_002839 Homo sapiens mitochondrial ribosomal protein S18C pseudogene 4 (MRPS1
 NG_002840 Homo sapiens mitochondrial ribosomal protein S21 pseudogene 8 (MRPS21
 NG_002841 Homo sapiens mitochondrial ribosomal protein L50 pseudogene 1 (MRPL501
 NG_002842 Homo sapiens mitochondrial ribosomal protein S31 pseudogene 1 (MRPS31
 NG_002843 Homo sapiens mitochondrial ribosomal protein L51 pseudogene 1 (MRPL511
 NG_002844 Homo sapiens mitochondrial ribosomal protein S23 pseudogene 1 (MRPS23
 NG_002845 Homo sapiens mitochondrial ribosomal protein 63 pseudogene 2 (MRP63P2;
 NG_002846 Homo sapiens mitochondrial ribosomal protein 63 pseudogene 3 (MRP63P3;
 NG_002847 Homo sapiens mitochondrial ribosomal protein 63 pseudogene 7 (MRP63P7;
 NG_002848 Homo sapiens mitochondrial ribosomal protein 63 pseudogene 8 (MRP63P8;
 NG_002849 Homo sapiens mitochondrial ribosomal protein 63 pseudogene 9 (MRP63P9;
 NG_002850 Homo sapiens mitochondrial ribosomal protein L11 pseudogene 3 (MRPL111
 NG_002851 Homo sapiens mitochondrial ribosomal protein L14 pseudogene 1 (MRPL141
 NG_002852 Homo sapiens mitochondrial ribosomal protein L15 pseudogene 1 (MRPL151
 NG_002853 Homo sapiens mitochondrial ribosomal protein L20 pseudogene 1 (MRPL201
 NG_002854 Homo sapiens mitochondrial ribosomal protein L22 pseudogene 1 (MRPL221
 NG_002855 Homo sapiens mitochondrial ribosomal protein L3 pseudogene 1 (MRPL3P1;
 NG_002856 Homo sapiens mitochondrial ribosomal protein L30 pseudogene 1 (MRPL301
 NG_002857 Homo sapiens mitochondrial ribosomal protein L32 pseudogene 1 (MRPL321
 NG_002858 Homo sapiens mitochondrial ribosomal protein L35 pseudogene 1 (MRPL351
 NG_002859 Homo sapiens mitochondrial ribosomal protein L35 pseudogene 2 (MRPL351
 NG_002860 Homo sapiens mitochondrial ribosomal protein L35 pseudogene 3 (MRPL351

NG_002861 Homo sapiens mitochondrial ribosomal protein L35 pseudogene 4 (MRPL35)
 NG_002862 Homo sapiens mitochondrial ribosomal protein L42 pseudogene 3 (MRPL42)
 NG_002863 Homo sapiens mitochondrial ribosomal protein L45 pseudogene 1 (MRPL45)
 NG_002864 Homo sapiens mitochondrial ribosomal protein L49 pseudogene 1 (MRPL49)
 NG_002865 Homo sapiens mitochondrial ribosomal protein L50 pseudogene 2 (MRPL50)
 NG_002866 Homo sapiens mitochondrial ribosomal protein L50 pseudogene 3 (MRPL50)
 NG_002867 Homo sapiens mitochondrial ribosomal protein L50 pseudogene 4 (MRPL50)
 NG_002868 Homo sapiens mitochondrial ribosomal protein L53 pseudogene 1 (MRPL53)
 NG_002869 Homo sapiens mitochondrial ribosomal protein S10 pseudogene 1 (MRPS10)
 NG_002870 Homo sapiens mitochondrial ribosomal protein S10 pseudogene 5 (MRPS10)
 NG_002871 Homo sapiens mitochondrial ribosomal protein S15 pseudogene 1 (MRPS15)
 NG_002872 Homo sapiens mitochondrial ribosomal protein S15 pseudogene 2 (MRPS15)
 NG_002873 Homo sapiens mitochondrial ribosomal protein S17 pseudogene 3 (MRPS17)
 NG_002874 Homo sapiens mitochondrial ribosomal protein S17 pseudogene 5 (MRPS17)
 NG_002875 Homo sapiens mitochondrial ribosomal protein S17 pseudogene 6 (MRPS17)
 NG_002876 Homo sapiens mitochondrial ribosomal protein S17 pseudogene 9 (MRPS17)
 NG_002877 Homo sapiens mitochondrial ribosomal protein S18A pseudogene 1 (MRPS1)
 NG_002878 Homo sapiens mitochondrial ribosomal protein S18B pseudogene 1 (MRPS1)
 NG_002879 Homo sapiens mitochondrial ribosomal protein S18B pseudogene 2 (MRPS1)
 NG_002880 Homo sapiens mitochondrial ribosomal protein S18C pseudogene 3 (MRPS1)
 NG_002881 Homo sapiens mitochondrial ribosomal protein S18C pseudogene 5 (MRPS1)
 NG_002882 Homo sapiens mitochondrial ribosomal protein S21 pseudogene 2 (MRPS21)
 NG_002883 Homo sapiens mitochondrial ribosomal protein S21 pseudogene 3 (MRPS21)
 NG_002884 Homo sapiens mitochondrial ribosomal protein S21 pseudogene 4 (MRPS21)
 NG_002885 Homo sapiens mitochondrial ribosomal protein S21 pseudogene 5 (MRPS21)
 NG_002886 Homo sapiens mitochondrial ribosomal protein S21 pseudogene 6 (MRPS21)
 NG_002887 Homo sapiens mitochondrial ribosomal protein S21 pseudogene 7 (MRPS21)
 NG_002888 Homo sapiens mitochondrial ribosomal protein S21 pseudogene 9 (MRPS21)
 NG_002889 Homo sapiens mitochondrial ribosomal protein S22 pseudogene 1 (MRPS22)
 NG_002890 Homo sapiens mitochondrial ribosomal protein S29 pseudogene 2 (MRPS29)
 NG_002891 Homo sapiens mitochondrial ribosomal protein S33 pseudogene 2 (MRPS33)
 NG_002892 Homo sapiens mitochondrial ribosomal protein S33 pseudogene 3 (MRPS33)
 NG_002893 Homo sapiens mitochondrial ribosomal protein S33 pseudogene 4 (MRPS33)
 NG_002894 Homo sapiens mitochondrial ribosomal protein S35 pseudogene 2 (MRPS35)
 NG_002895 Homo sapiens mitochondrial ribosomal protein S35 pseudogene 3 (MRPS35)
 NG_002896 Homo sapiens mitochondrial ribosomal protein S5 pseudogene 1 (MRPS5P1)
 NG_002897 Homo sapiens mitochondrial ribosomal protein S5 pseudogene 2 (MRPS5P2)
 NG_002898 Homo sapiens mitochondrial ribosomal protein S6 pseudogene 4 (MRPS6P4)
 NG_002899 Homo sapiens mitochondrial ribosomal protein S7 pseudogene 1 (MRPS7P1)
 NG_002900 Homo sapiens mitochondrial ribosomal protein S5 pseudogene (MRPS5P4)
 NG_002901 Homo sapiens PCNA pseudogene pFPCNA (LOC359805) on chromosome
 NG_002902 Homo sapiens PCNA pseudogene p1PCNA (LOC359806) on chromosome 4
 NG_002903 Homo sapiens mitochondrial ribosomal protein S10 pseudogene (MRPS10P;
 NG_002904 Homo sapiens mitochondrial ribosomal protein S29 pseudogene 1 (MRPS29
 NG_002905 Homo sapiens chloride channel, nucleotide-sensitive, 1B (CLNS1B) pseudog
 NG_002906 Homo sapiens VENT-like homeobox 2 pseudogene 4 (VENTX2P4) on chrom
 NG_002907 Homo sapiens mitochondrial ribosomal protein L42 pseudogene 2 (MRPL42)
 NG_002908 Homo sapiens mitochondrial ribosomal protein L42 pseudogene 4 (MRPL42)
 NG_002909 Homo sapiens VENT-like homeobox 2 pseudogene 2 (VENTX2P2) on chrom
 NG_002910 Homo sapiens mitochondrial ribosomal protein L10 pseudogene (LOC34895;
 NG_002911 Homo sapiens VENT-like homeobox 2 pseudogene 3 (VENTX2P3) on chrom
 NG_002912 Homo sapiens mitochondrial ribosomal protein L30 pseudogene 2 (MRPL30)
 NG_002913 Homo sapiens mitochondrial ribosomal protein L39 pseudogene (LOC35981;
 NG_002914 Homo sapiens mitochondrial ribosomal protein L42 pseudogene 5 (MRPL42)
 NG_002915 Homo sapiens peroxiredoxin 2 pseudogene 1 (PRDX2P1) on chromosome 1
 NG_002916 Homo sapiens MHC class I polypeptide-related sequence E (MICE) pseudog
 NG_002917 Homo sapiens PAI-1 mRNA-binding protein pseudogene (LOC359996) on chr

NG_002918 Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex, su
 NG_002919 Homo sapiens CHRNA7 (cholinergic receptor, nicotinic, alpha polypeptide 7,
 NG_002920 Homo sapiens CHRNA7 (cholinergic receptor, nicotinic, alpha polypeptide 7,
 NG_002921 Homo sapiens BCL6 co-repressor pseudogene (LOC360000) on chromosom
 NG_002922 Homo sapiens interferon alpha-L pseudogene (G13P1) on chromosome 9
 NG_002923 Homo sapiens glyceraldehyde-3-phosphate dehydrogenase pseudogene (LC
 NG_002924 Homo sapiens glyceraldehyde-3-phosphate dehydrogenase pseudogene (LC
 NG_002926 Homo sapiens cytochrome c, somatic pseudogene (HCP46) on chromosom
 NG_002927 Homo sapiens cytochrome c, somatic pseudogene (LOC360009) on chromo
 NG_002928 Homo sapiens RAD17 homolog (S. pombe) pseudogene 2 (RAD17P2) on ch
 NG_002929 Homo sapiens RAD17 homolog (S. pombe) pseudogene 1 (RAD17P1) on ch
 NG_002930 Homo sapiens TAF9 pseudogene 1 (TAF9P1) on chromosome Y
 NG_002931 Homo sapiens TAF9 pseudogene 2 (TAF9P2) on chromosome Y
 NG_002932 Homo sapiens discs, large homolog 7 (Drosophila) pseudogene (LOC36001
 NG_002933 Homo sapiens raft-linking protein pseudogene (LOC360015) on chromosom
 NG_002934 Homo sapiens capicua homolog (Drosophila) pseudogene (LOC360016) on
 NG_002935 Homo sapiens capicua homolog (Drosophila) pseudogene (LOC360017) on
 NG_002936 Homo sapiens HbxAg transactivated protein 2 pseudogene (LOC360018) on
 NG_002937 Homo sapiens keratin 18 pseudogene 10 (KRT18P10) on chromosome Y
 NG_002938 Homo sapiens PC4 and SFRS1 interacting protein 2 pseudogene (LOC3600
 NG_002939 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 12B pset
 NG_002940 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 12B (LOC
 NG_002941 Homo sapiens ribosomal protein L41 pseudogene (LOC286568) on chromos
 NG_002942 Homo sapiens ribosomal protein L41 pseudogene (LOC286570) on chromos
 NG_002943 Homo sapiens Intersectin 2 pseudogene (LOC360027) on chromosome Y
 NG_002944 Homo sapiens Intersectin 2 pseudogene (LOC360026) on chromosome Y
 NG_002945 Homo sapiens Intersectin 2 pseudogene (LOC360025) on chromosome Y
 NG_002946 Homo sapiens Intersectin 2 pseudogene (LOC360024) on chromosome Y
 NG_002947 Homo sapiens tubulin, beta polypeptide 4, member Q pseudogene (LOC1401
 NG_002948 Homo sapiens tubulin, beta polypeptide 4, member Q pseudogene (LOC349
 NG_002949 Homo sapiens hypothetical protein MGC23909 pseudogene (LOC360028) on
 NG_002950 Homo sapiens hypothetical protein MGC45134 (MGC45134) pseudogene on
 NG_002951 Homo sapiens cyclic-AMP-dependent transcription factor ATF-4 pseudogene
 NG_002952 Homo sapiens hypothetical protein BC016683 pseudogene (LOC360029) on
 NG_002953 Homo sapiens cytochrome c, somatic pseudogene (HCP1) on chromosome
 NG_002954 Homo sapiens cytochrome c, somatic pseudogene (HCP3) on chromosome
 NG_002955 Homo sapiens cytochrome c, somatic pseudogene (HCP4) on chromosome
 NG_002956 Homo sapiens cytochrome c, somatic pseudogene (HCP5) on chromosome
 NG_002957 Homo sapiens cytochrome c, somatic pseudogene (HCP6) on chromosome
 NG_002958 Homo sapiens cytochrome c, somatic pseudogene (HCP7) on chromosome
 NG_002959 Homo sapiens cytochrome c, somatic pseudogene (HCP8) on chromosome
 NG_002960 Homo sapiens cytochrome c, somatic pseudogene (HCP9) on chromosome
 NG_002961 Homo sapiens cytochrome c, somatic pseudogene (HCP10) on chromosome
 NG_002962 Homo sapiens cytochrome c, somatic pseudogene (HCP11) on chromosome
 NG_002963 Homo sapiens cytochrome c, somatic pseudogene (HCP12) on chromosome
 NG_002964 Homo sapiens cytochrome c, somatic pseudogene (HCP13) on chromosome
 NG_002965 Homo sapiens cytochrome c, somatic pseudogene (HCP14) on chromosome
 NG_002966 Homo sapiens cytochrome c, somatic pseudogene (HCP16) on chromosome
 NG_002967 Homo sapiens cytochrome c, somatic pseudogene (HCP17) on chromosome
 NG_002968 Homo sapiens cytochrome c, somatic pseudogene (HCP18) on chromosome
 NG_002969 Homo sapiens cytochrome c, somatic pseudogene (HCP19) on chromosome
 NG_002970 Homo sapiens cytochrome c, somatic pseudogene (HCP20) on chromosome
 NG_002971 Homo sapiens cytochrome c, somatic pseudogene (HCP21) on chromosome
 NG_002972 Homo sapiens cytochrome c, somatic pseudogene (HCP22) on chromosome
 NG_002973 Homo sapiens cytochrome c, somatic pseudogene (HCP23) on chromosome
 NG_002974 Homo sapiens cytochrome c, somatic pseudogene (HCP24) on chromosome
 NG_002975 Homo sapiens cytochrome c, somatic pseudogene (HCP25) on chromosome

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NG_002976 Homo sapiens cytochrome c, somatic pseudogene (HCP26) on chromosome
 NG_002977 Homo sapiens cytochrome c, somatic pseudogene (HCP27) on chromosome
 NG_002978 Homo sapiens cytochrome c, somatic pseudogene (HCP28) on chromosome
 NG_002979 Homo sapiens cytochrome c, somatic pseudogene (HCP29) on chromosome
 NG_002980 Homo sapiens cytochrome c, somatic pseudogene (HCP30) on chromosome
 NG_002981 Homo sapiens cytochrome c, somatic pseudogene (HCP32) on chromosome
 NG_002982 Homo sapiens cytochrome c, somatic pseudogene (HCP33) on chromosome
 NG_002983 Homo sapiens cytochrome c, somatic pseudogene (HCP34) on chromosome
 NG_002984 Homo sapiens cytochrome c, somatic pseudogene (HCP35) on chromosome
 NG_002985 Homo sapiens cytochrome c, somatic pseudogene 1 (CYCSP1) on chromos
 NG_002986 Homo sapiens cytochrome c, somatic pseudogene (HCP37) on chromosome
 NG_002987 Homo sapiens cytochrome c, somatic pseudogene (HCP38) on chromosome
 NG_002988 Homo sapiens cytochrome c, somatic pseudogene (HCP39) on chromosome
 NG_002989 Homo sapiens cytochrome c, somatic pseudogene (HCP40) on chromosome
 NG_002990 Homo sapiens cytochrome c, somatic pseudogene (HCP41) on chromosome
 NG_002991 Homo sapiens cytochrome c, somatic pseudogene (HCP42) on chromosome
 NG_002992 Homo sapiens cytochrome c, somatic pseudogene (HCP43) on chromosome
 NG_002993 Homo sapiens cytochrome c, somatic pseudogene (HCP44) on chromosome
 NG_002994 Homo sapiens cytochrome c, somatic pseudogene (HCP45) on chromosome
 NG_002995 Homo sapiens cytochrome c, somatic pseudogene (HCP48) on chromosome
 NG_002996 Homo sapiens thymosin-like 5 (TMSL5) pseudogene on chromosome 11
 NG_002997 Homo sapiens thymosin-like 7 (TMSL7) pseudogene on chromosome X
 NG_002998 Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 4, 9
 NG_002999 Homo sapiens cytochrome c, somatic pseudogene (HCP49) on chromosome
 NG_003000 Homo sapiens RNA binding motif protein, Y-linked, family 1, member H (RBM
 NG_003008 Homo sapiens nuclease sensitive element binding protein 1 pseudogene (bA
 NG_003009 Homo sapiens argininosuccinate synthetase pseudogene 8 (ASSP8) on chr
 NG_003010 Homo sapiens cytochrome c oxidase subunit Vb-like 1 (COX5BL1) pseudoge
 NG_003011 Homo sapiens glyceraldehyde-3-phosphate dehydrogenase-like 4 (GAPDL4)
 NG_003012 Homo sapiens glycine dehydrogenase (decarboxylase) pseudogene (GLDCF
 NG_003013 Homo sapiens general transcription factor IIF, polypeptide 2-like (GTF2F2L)
 NG_003014 Homo sapiens heat shock 90kDa protein 1, alpha-like 2 (HSPCAL2) pseudog
 NG_003015 Homo sapiens heat shock 90kDa protein 1, beta pseudogene 1 (HSPCP1) o
 NG_003016 Homo sapiens lactate dehydrogenase A-like 1 (LDHAL1) pseudogene on chr
 NG_003017 Homo sapiens sorcin-like (SRIL) pseudogene on chromosome 4
 NG_003018 Homo sapiens v-ras-1 murine leukemia viral oncogene homolog 1 pseudog
 NG_003019 Homo sapiens actin, beta pseudogene 2 (ACTBP2) on chromosome 5
 NG_003020 Homo sapiens actin, beta pseudogene 4 (ACTBP4) on chromosome 5
 NG_003021 Homo sapiens argininosuccinate synthetase pseudogene 10 (ASSP10) on cl
 NG_003022 Homo sapiens argininosuccinate synthetase pseudogene 9 (ASSP9) on chr
 NG_003023 Homo sapiens OFD1 pseudogene 1 (OFD1P1) on chromosome 5
 NG_003024 Homo sapiens diazepam binding inhibitor-like 1 (DBIL1) pseudogene on chr
 NG_003025 Homo sapiens chemokine ligand 14, chemokine ligand 15 transcription unit (
 NG_003026 Homo sapiens endogenous retroviral pol gene-like sequence 2 (ERPL2) pse
 NG_003027 Homo sapiens glyceraldehyde-3-phosphate dehydrogenase-like 16 (GAPDL'
 NG_003028 Homo sapiens ferritin, heavy polypeptide-like 10 (FTHL10) pseudogene on c
 NG_003029 Homo sapiens gap junction protein, alpha 1, 43kDa (connexin 43) pseudoge
 NG_003030 Homo sapiens glutamate-ammonia ligase (glutamine synthase)-like 1 (GLUL
 NG_003031 Homo sapiens hypoxanthine phosphoribosyltransferase pseudogene 2 (HPR
 NG_003032 Homo sapiens moesin-like 1 (MSN1) pseudogene on chromosome 5
 NG_003033 Homo sapiens ribosomal protein S17 pseudogene 2 (RPS17P2) on chromos
 NG_003034 Homo sapiens ribosomal protein S20 pseudogene 3 (RPS20P3) on chromos
 NG_003035 Homo sapiens ribosomal protein S20 pseudogene 4 (RPS20P4) on chromos
 NG_003036 Homo sapiens t-complex 1-like 2 (TCP1L2) pseudogene on chromosome 5
 NG_003037 Homo sapiens X-box binding protein pseudogene 1 (XBPP1) on chromosome
 NG_003038 Homo sapiens eukaryotic translation elongation factor 1 beta 3 (EEF1B3) ps
 NG_003039 Homo sapiens actin, gamma pseudogene 10 (ACTGP10) on chromosome X

NG_003040 Homo sapiens brain cytoplasmic RNA 1, pseudogene 1 (BCYRN1P1) on chrn
 NG_003041 Homo sapiens RNA binding motif protein, Y-linked, family 2, member B pseu
 NG_003042 Homo sapiens RNA binding motif protein, Y-linked, family 2, member C pseu
 NG_003043 Homo sapiens RNA binding motif protein, Y-linked, family 2, member D pseu
 NG_003044 Homo sapiens centromere protein C2, 140kDa (CENPC2) pseudogene on ch
 NG_003045 Homo sapiens eukaryotic translation elongation factor 1 beta 1 (EEF1B1) psi
 NG_003061 Homo sapiens DNA segment on chromosome 6 (unique, pseudogene) 2723
 NG_003062 Homo sapiens ribosomal protein L7a pseudogene 1 (RPL7AP1) on chromos
 NG_003063 Homo sapiens zinc finger protein 381, Y-linked pseudogene (ZNF381P) on c
 NG_003064 Homo sapiens RNA binding motif protein, Y-linked, family 2, member A pseu
 NG_003070 Homo sapiens RNA binding motif protein, Y-linked, family 2, pseudogene (LC
 NG_003072 Homo sapiens MADS box transcription enhancer factor 2, polypeptide A pse
 NG_003073 Homo sapiens protein kinase, cAMP-dependent, regulatory, type I, alpha pse
 NG_003074 Homo sapiens RNA binding motif protein, Y-linked, family 2, pseudogene (LC
 NG_003075 Homo sapiens RNA binding motif protein, Y-linked, family 2, pseudogene (LC
 NG_003076 Homo sapiens ring finger protein 134 pseudogene 1 (RNF134P1) on chromo
 NG_003077 Homo sapiens testis specific protein, Y-linked pseudogene 1 (TSPYP1) on cl
 NG_003078 Homo sapiens testis specific protein, Y-linked pseudogene 2 (TSPYP2) on cl
 NG_003079 Homo sapiens testis specific protein, Y-linked pseudogene 3 (TSPYP3) on cl
 NG_003080 Homo sapiens testis specific protein, Y-linked pseudogene 4 (TSPYP4) on cl
 NG_003081 Homo sapiens RNA binding motif protein, Y-linked, family 2, pseudogene (LC
 NG_003082 Homo sapiens RNA binding motif protein, Y-linked, family 2, pseudogene (LC
 NG_003083 Homo sapiens RNA binding motif protein, Y-linked, family 2, pseudogene (LC
 NG_003084 Homo sapiens RNA binding motif protein, Y-linked, family 2, pseudogene (LC
 NG_003085 Homo sapiens RNA binding motif protein, Y-linked, family 2, pseudogene (LC
 NG_003086 Homo sapiens RNA binding motif protein, Y-linked, family 2, pseudogene (LC
 NG_003087 Homo sapiens RNA binding motif protein, Y-linked, family 2, pseudogene (LC
 NG_003088 Homo sapiens RNA binding motif protein, Y-linked, family 2, pseudogene (LC
 NG_003089 Homo sapiens RNA binding motif protein, Y-linked, family 2, pseudogene (LC
 NG_003090 Homo sapiens RNA binding motif protein, Y-linked, family 2, pseudogene (LC
 NG_003091 Homo sapiens RNA binding motif protein, Y-linked, family 2, pseudogene (LC
 NG_003092 Homo sapiens RNA binding motif protein, Y-linked, family 2, pseudogene (LC
 NG_003093 Homo sapiens testis specific protein, Y-linked pseudogene 5 (TSPYP5) on cl
 NG_003094 Homo sapiens X Kell blood group precursor-related, Y-linked pseudogene 1
 NG_003095 Homo sapiens X Kell blood group precursor-related, Y-linked pseudogene 2
 NG_003096 Homo sapiens X Kell blood group precursor-related, Y-linked pseudogene 3
 NG_003097 Homo sapiens X Kell blood group precursor-related, Y-linked pseudogene 4
 NG_003098 Homo sapiens X Kell blood group precursor-related, Y-linked pseudogene 5
 NG_003099 Homo sapiens X Kell blood group precursor-related, Y-linked pseudogene (X
 NG_003100 Homo sapiens choline kinase-like, carnitine palmitoyltransferase 1B (muscle)
 NG_003101 Homo sapiens ubiquitin-conjugating enzyme E2L 1 (UBE2L1) pseudogene o
 NG_003102 Homo sapiens COP9 constitutive photomorphogenic homolog subunit 5 pse
 NG_003103 Homo sapiens COP9 pseudogene (LOC375350) on chromosome 3
 NG_003104 Homo sapiens zinc finger protein 91 homolog (mouse), ciliary neurotrophic fe
 NG_003105 Homo sapiens E2F transcription factor 6 pseudogene (LOC376818) on chr
 NG_003106 Homo sapiens E2F transcription factor 6 pseudogene (LOC386610) on chr
 NG_003107 Homo sapiens solute carrier family 25 (mitochondrial carrier; adenine nucle
 NG_003108 Homo sapiens similar to v-raf murine sarcoma viral oncogene homolog B1 ps
 NG_003109 Homo sapiens osteoclast stimulating factor 1 pseudogene (OSTF1P) on chr
 NG_003110 Homo sapiens LAG1 longevity assurance homolog 1 (S. cerevisiae), growth
 NG_003111 Homo sapiens serine/threonine kinase 6-like pseudogene (STK6LP) on chro
 NG_003114 Homo sapiens OFD1 pseudogene Y-linked 1 (OFDYP1) on chromosome Y
 NG_003115 Homo sapiens OFD1 pseudogene Y-linked 2 (OFDYP2) on chromosome Y
 NG_003116 Homo sapiens OFD1 pseudogene Y-linked 3 (OFDYP3) on chromosome Y
 NG_003117 Homo sapiens OFD1 pseudogene Y-linked 4 (OFDYP4) on chromosome Y
 NG_003118 Homo sapiens OFD1 pseudogene Y-linked 5 (OFDYP5) on chromosome Y
 NG_003119 Homo sapiens OFD1 pseudogene Y-linked 6 (OFDYP6) on chromosome Y

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NG_003120 Homo sapiens OFD1 pseudogene Y-linked 7 (OFDYP7) on chromosome Y
 NG_003121 Homo sapiens OFD1 pseudogene Y-linked 8 (OFDYP8) on chromosome Y
 NG_003122 Homo sapiens OFD1 pseudogene Y-linked 9 (OFDYP9) on chromosome Y
 NG_003125 Homo sapiens OFD1 pseudogene Y-linked 11 (OFDYP11) on chromosome Y
 NG_003126 Homo sapiens OFD1 pseudogene Y-linked 12 (OFDYP12) on chromosome Y
 NG_003127 Homo sapiens OFD1 pseudogene Y-linked 13 (OFDYP13) on chromosome Y
 NG_003128 Homo sapiens OFD1 pseudogene Y-linked 14 (OFDYP14) on chromosome Y
 NG_003129 Homo sapiens OFD1 pseudogene Y-linked 15 (OFDYP15) on chromosome Y
 NG_003130 Homo sapiens chromodomain protein, Y-linked 10 pseudogene (CDY10P) or
 NG_003131 Homo sapiens chromodomain protein, Y-linked 3 pseudogene (CDY3P) on c
 NG_003132 Homo sapiens chromodomain protein, Y-linked 5 pseudogene (CDY5P) on c
 NG_003133 Homo sapiens chromodomain protein, Y-linked 9 pseudogene (CDY9P) on c
 NG_003134 Homo sapiens family with sequence similarity 8, member A7 pseudogene (F
 NG_003135 Homo sapiens family with sequence similarity 8, member A8 pseudogene (F
 NG_003136 Homo sapiens family with sequence similarity 8, member A9 pseudogene (F
 NG_003137 Homo sapiens family with sequence similarity 8, member A1 pseudogene (L
 NG_003138 Homo sapiens chromodomain protein, Y-linked 4 pseudogene (CDY4P) on c
 NG_003139 Homo sapiens chromodomain protein, Y-linked 6 pseudogene (CDY6P) on c
 NG_003140 Homo sapiens chromodomain protein, Y-linked 7 pseudogene (CDY7P) on c
 NG_003141 Homo sapiens chromodomain protein, Y-linked 8 pseudogene (CDY8P) on c
 NG_003142 Homo sapiens chromodomain protein, Y-linked 12 pseudogene (CDY12P) or
 NG_003143 Homo sapiens chromodomain protein, Y-linked 13 pseudogene (CDY13P) or
 NG_003144 Homo sapiens chromodomain protein, Y-linked 14 pseudogene (CDY14P) or
 NG_003145 Homo sapiens chromodomain protein, Y-linked 15 pseudogene (CDY15P) or
 NG_003146 Homo sapiens chromodomain protein, Y-linked 16 pseudogene (CDY16P) or
 NG_003147 Homo sapiens chromodomain protein, Y-linked 17 pseudogene (CDY17P) or
 NG_003148 Homo sapiens chromodomain protein, Y-linked 18 pseudogene (CDY18P) or
 NG_003149 Homo sapiens chromodomain protein, Y-linked 19 pseudogene (CDY19P) or
 NG_003150 Homo sapiens chromodomain protein, Y-linked 20 pseudogene (CDY20P) or
 NG_003151 Homo sapiens chromodomain protein, Y-linked 21 pseudogene (CDY21P) or
 NG_003152 Homo sapiens chromodomain protein, Y-linked 22 pseudogene (CDY22P) or
 NG_003153 Homo sapiens chromodomain protein, Y-linked 23 pseudogene (CDY23P) or
 NG_003154 Homo sapiens chromodomain protein, Y-linked 11 pseudogene (CDY11P) or
 NG_003155 Homo sapiens ribosomal protein L29 pseudogene 1 (RPL29P1) on chromos
 NG_003156 Homo sapiens ribosomal protein L39 pseudogene 3 (RPL39P3) on chromos
 NG_003157 Homo sapiens v-raf murine sarcoma 3B11 viral oncogene homolog pseudoge
 NG_003158 Homo sapiens telomeric repeat binding factor (NIMA-interacting) 1 pseudoge
 NG_003159 Homo sapiens pseudogene of CXYorf1 (CXYorf1P) on chromosome 18
 NG_003160 Homo sapiens ribosomal protein L24 pseudogene 4 (RPL24P4) on chromos
 NG_003162 Homo sapiens actin, beta pseudogene 9 (ACTBP9) on chromosome 18
 NG_003163 Homo sapiens creatine kinase B pseudogene 1 (CKBP1) on chromosome 1E
 NG_003164 Homo sapiens transferrin pseudogene (TFP) on chromosome 3
 NG_003165 Homo sapiens platelet-activating factor acetylhydrolase, isoform lb, pseudog
 NG_003166 Homo sapiens olfactory receptor, family 7, subfamily A, member 8 pseudoge
 NG_003167 Homo sapiens platelet-activating factor acetylhydrolase, isoform lb, pseudog
 NG_003168 Homo sapiens ubiquitin specific protease 9, Y-linked pseudogene (LOC3873
 NG_003169 Homo sapiens ubiquitin specific protease 9, Y-linked pseudogene (LOC3873
 NG_003170 Homo sapiens ubiquitin specific protease 9, Y-linked pseudogene (LOC3873
 NG_003171 Homo sapiens USP9Y pseudogene 1 (USP9YP1) on chromosome Y
 NG_003172 Homo sapiens USP9Y pseudogene 2 (USP9YP2) on chromosome Y
 NG_003173 Homo sapiens ubiquitin specific protease 9, Y-linked pseudogene (LOC3873
 NG_003180 Homo sapiens cytochrome P450, family 2, subfamily D, polypeptide 6 genot
 NG_003183 Homo sapiens BRCA1 pseudogene (LBRCA1) on chromosome 17
 NG_003186 Homo sapiens transcription elongation factor A (SII), 1 pseudogene (LOC39
 NG_003187 Homo sapiens deafness dystonia pseudogene (DDPP) on chromosome 2
 NG_003188 Homo sapiens chemokine (C-C motif) ligand 3-like 2 (CCL3L2) pseudogene
 NG_003189 Homo sapiens ribosomal protein S26 pseudogene 6 (RPS26P6) on chromos

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NG_003190 Homo sapiens pseudogene of ribosomal protein L9 (LOC388147) on chromo
 NG_003193 Homo sapiens PR/SET domain containing protein 07 pseudogene (SET07p)
 NG_003194 Homo sapiens olfactory receptor, family 2, subfamily B, member 8 pseudoge
 NG_003195 Homo sapiens olfactory receptor, family 52, subfamily L, member 2 pseudog
 NG_003196 Homo sapiens olfactory receptor, family 10, subfamily D, member 4 pseudog
 NG_003197 Homo sapiens olfactory receptor, family 52, subfamily E, member 1 pseudog
 NG_003198 Homo sapiens olfactory receptor, family 4, subfamily K, member 3 pseudoge
 NG_003200 Homo sapiens ELK2, member of ETS oncogene family, pseudogene 2 (ELK2
 NG_003201 Homo sapiens olfactory receptor, family 4, subfamily G, member 2 pseudoge
 NG_003215 Homo sapiens eukaryotic translation initiation factor 2, subunit 3 gamma, 52
 NG_003216 Homo sapiens ribosomal protein S25 pseudogene (LOC283114) on chromos
 NG_003217 Homo sapiens heat shock 60kDa protein 1 pseudogene (LOC283320) on chi
 NG_003218 Homo sapiens synaptogyrin 2 pseudogene (LOC283698) on chromosome 1E
 NG_003219 Homo sapiens transcription elongation factor B (SII), polypeptide 1 pseudog
 NG_003221 Homo sapiens olfactory receptor, family 11, subfamily I, member 1 pseudoge
 NG_003222 Homo sapiens olfactory receptor, family 4, subfamily G, member 3 pseudoge
 NG_003230 Homo sapiens olfactory receptor, family 5, subfamily D, member 11 pseudog
 NG_003253 Homo sapiens cytochrome c oxidase, subunit 8B pseudogene (COX8B) on c
 NG_003254 Homo sapiens immunoglobulin heavy constant epsilon P2 (IGHEP2) pseudo
 NG_003255 Homo sapiens Rhesus blood group cluster (RHD/RHCE) on chromosome
 NG_003256 Homo sapiens gonadotropin-releasing hormone receptor 2 pseudogene (GN
 NG_003258 Homo sapiens folate hydrolase 2 (FOLH2) pseudogene on chromosome 11
 NG_003259 Homo sapiens ribosomal protein S9 pseudogene 1 (RPS9P1) on chromoso
 NG_004075 Homo sapiens laminin receptor 1 pseudogene 14 (LAMR1P14) on chromoso
 NG_004077 Homo sapiens thiopurine S-methyltransferase pseudogene (LOC40650) on
 NG_004085 Homo sapiens protein tyrosine phosphatase type IVA pseudogene 1 (PTP4A
 NG_004086 Homo sapiens olfactory receptor, family 5, subfamily J, member 1 pseudoge
 NG_004087 Homo sapiens olfactory receptor, family 7, subfamily A, member 2 pseudoge
 NG_004088 Homo sapiens olfactory receptor, family 2, subfamily E, member 1 pseudoge
 NG_004089 Homo sapiens olfactory receptor, family 4, subfamily H, member 12 pseudog
 NG_004091 Homo sapiens mitochondrial ribosomal protein S17 pseudogene (MRPS17P
 NG_004092 Homo sapiens mitochondrial ribosomal protein S17 pseudogene (MRPS17P
 NG_004093 Homo sapiens Nanog homeobox pseudogene 8 (NANOGP8) on chromosom
 NG_004095 Homo sapiens Nanog homeobox pseudogene 3 (NANOGP3) on chromosom
 NG_004096 Homo sapiens Nanog homeobox pseudogene 10 (NANOGP10) on chromosom
 NG_004097 Homo sapiens Nanog homeobox pseudogene 9 (NANOGP9) on chromosom
 NG_004098 Homo sapiens NANOG homeobox pseudogene 7 (NANOGP7) on chromosom
 NG_004099 Homo sapiens NANOG homeobox pseudogene 2 (NANOGP2) on chromosom
 NG_004100 Homo sapiens NANOG homeobox pseudogene 4 (NANOGP4) on chromosom
 NG_004101 Homo sapiens NANOG homeobox pseudogene 5 (NANOGP5) on chromosom
 NG_004102 Homo sapiens NANOG pseudogene 6 (NANOGP6) on chromosome 10
 NG_004103 Homo sapiens NANOG homeobox pseudogene 11 (NANOGP11) on chromo
 NG_004109 Homo sapiens YTH domain family 2 pseudogene (YTHDF2P) on chromosom
 NG_004110 Homo sapiens ribonuclease H1 pseudogene 2 (RNASEH1P2) on chromosom
 NG_004111 Homo sapiens ribonuclease H1 pseudogene 1 (RNASEH1P1) on chromosom
 NG_004112 Homo sapiens endogenous retroviral family W, env(C7), member 1 (synofin
 NG_004113 Homo sapiens CCL3L1-CCL4L1 chemokine gene cluster (CCL3L1-CCL4L1)
 NG_004115 Homo sapiens VKORC1 pseudogene 1 (LOC414355) on chromosome X
 NG_004116 Homo sapiens VKORC1 pseudogene 2 (LOC414357) on chromosome 1
 NG_004122 Homo sapiens olfactory receptor, family 7, subfamily E, member 22 pseudog
 NG_004123 Homo sapiens olfactory receptor, family 7, subfamily E, member 21 pseudog
 NG_004124 Homo sapiens olfactory receptor, family 5, subfamily E, member 1 pseudoge
 NG_004125 Homo sapiens olfactory receptor, family 10, subfamily D, member 3 pseudog
 NG_004126 Homo sapiens olfactory receptor, family 10, subfamily D, member 1 pseudog
 NG_004127 Homo sapiens olfactory receptor, family 7, subfamily E, member 66 pseudog
 NG_004128 Homo sapiens olfactory receptor, family 7, subfamily E, member 47 pseudog
 NG_004129 Homo sapiens olfactory receptor, family 7, subfamily E, member 36 pseudog

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NG_004245 Homo sapiens olfactory receptor, family 10, subfamily Y, member 1 pseudog
NG_004246 Homo sapiens olfactory receptor, family 10, subfamily W, member 1 (OR10V
NG_004247 Homo sapiens olfactory receptor, family 10, subfamily V, member 3 pseudog
NG_004248 Homo sapiens olfactory receptor, family 10, subfamily V, member 2 pseudog
NG_004249 Homo sapiens olfactory receptor, family 10, subfamily Q, member 2 pseudog
NG_004250 Homo sapiens olfactory receptor, family 7, subfamily M, member 1 pseudoge
NG_004251 Homo sapiens olfactory receptor, family 7, subfamily E, member 110 pseudo
NG_004252 Homo sapiens olfactory receptor, family 6, subfamily L, member 2 pseudoge
NG_004253 Homo sapiens olfactory receptor, family 7, subfamily E, member 116 pseudo
NG_004254 Homo sapiens olfactory receptor, family 7, subfamily E, member 108 pseudo
NG_004255 Homo sapiens olfactory receptor, family 2, subfamily AM, member 1 pseudog
NG_004256 Homo sapiens olfactory receptor, family 7, subfamily A, member 125 pseudo
NG_004257 Homo sapiens olfactory receptor, family 9, subfamily P, member 1 pseudoge
NG_004258 Homo sapiens olfactory receptor, family 9, subfamily N, member 1 pseudoge
NG_004259 Homo sapiens olfactory receptor, family 4, subfamily F, member 7 pseudoge
NG_004260 Homo sapiens olfactory receptor, family 2, subfamily W, member 6 pseudoge
NG_004261 Homo sapiens olfactory receptor, family 2, subfamily W, member 4 pseudoge
NG_004262 Homo sapiens olfactory receptor, family 7, subfamily E, member 99 pseudog
NG_004263 Homo sapiens olfactory receptor, family 5, subfamily M, member 14 pseudog
NG_004264 Homo sapiens olfactory receptor, family 7, subfamily A, member 129 pseudo
NG_004265 Homo sapiens olfactory receptor, family 7, subfamily A, member 122 pseudo
NG_004266 Homo sapiens olfactory receptor, family 7, subfamily E, member 100 pseudo
NG_004267 Homo sapiens olfactory receptor, family 5, subfamily AC, member 1 pseudog
NG_004268 Homo sapiens olfactory receptor, family 4, subfamily G, member 6 pseudoge
NG_004269 Homo sapiens olfactory receptor, family 9, subfamily H, member 1 pseudoge
NG_004270 Homo sapiens olfactory receptor, family 6, subfamily R, member 1 pseudoge
NG_004271 Homo sapiens olfactory receptor, family 6, subfamily K, member 1 pseudoge
NG_004272 Homo sapiens olfactory receptor, family 2, subfamily T, member 7 (OR2T7) f
NG_004273 Homo sapiens olfactory receptor, family 2, subfamily L, member 9 pseudoge
NG_004274 Homo sapiens olfactory receptor, family 2, subfamily L, member 6 pseudoge
NG_004275 Homo sapiens olfactory receptor, family 2, subfamily L, member 5 (OR2L5) f
NG_004276 Homo sapiens olfactory receptor, family 2, subfamily AQ, member 1 pseudog
NG_004277 Homo sapiens olfactory receptor, family 10, subfamily AE, member 1 pseudo
NG_004278 Homo sapiens olfactory receptor, family 10, subfamily AA, member 1 pseudo
NG_004279 Homo sapiens olfactory receptor, family 2, subfamily W, member 2 pseudoge
NG_004280 Homo sapiens olfactory receptor, family 2, subfamily B, member 7 pseudoge
NG_004281 Homo sapiens olfactory receptor, family 51, subfamily N, member 1 pseudog
NG_004282 Homo sapiens olfactory receptor, family 52, subfamily P, member 2 pseudog
NG_004283 Homo sapiens olfactory receptor, family 2, subfamily AP, member 1 (OR2AP
NG_004284 Homo sapiens olfactory receptor, family 11, subfamily M, member 1 pseudog
NG_004285 Homo sapiens olfactory receptor, family 9, subfamily K, member 1 pseudoge
NG_004286 Homo sapiens olfactory receptor, family 5, subfamily AV, member 1 pseudog
NG_004287 Homo sapiens olfactory receptor, family 6, subfamily P, member 1 (OR6P1) f
NG_004288 Homo sapiens olfactory receptor, family 6, subfamily K, member 4 pseudoge
NG_004289 Homo sapiens olfactory receptor, family 2, subfamily AI, member 1 pseudoge
NG_004290 Homo sapiens olfactory receptor, family 2, subfamily A, member 15 pseudog
NG_004291 Homo sapiens olfactory receptor, family 7, subfamily E, member 136 pseudo
NG_004292 Homo sapiens olfactory receptor, family 2, subfamily A, member 3 pseudoge
NG_004293 Homo sapiens olfactory receptor, family 6, subfamily D, member 1 pseudoge
NG_004294 Homo sapiens olfactory receptor, family 7, subfamily E, member 39 pseudog
NG_004295 Homo sapiens olfactory receptor, family 4, subfamily C, member 50 pseudog
NG_004296 Homo sapiens olfactory receptor, family 4, subfamily D, member 12 pseudog
NG_004297 Homo sapiens olfactory receptor, family 7, subfamily E, member 149 pseudo
NG_004298 Homo sapiens olfactory receptor, family 10, subfamily AF, member 1 pseudo
NG_004299 Homo sapiens olfactory receptor, family 8, subfamily V, member 1 pseudoge
NG_004300 Homo sapiens olfactory receptor, family 5, subfamily BT, member 1 pseudog
NG_004301 Homo sapiens olfactory receptor, family 8, subfamily T, member 1 pseudoge

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NG_004302 Homo sapiens olfactory receptor, family 11, subfamily P, member 1 pseudog
NG_004303 Homo sapiens olfactory receptor, family 7, subfamily E, member 148 pseudo
NG_004304 Homo sapiens olfactory receptor, family 52, subfamily Z, member 1 pseudog
NG_004305 Homo sapiens olfactory receptor, family 52, subfamily M, member 2 pseudog
NG_004307 Homo sapiens olfactory receptor, family 7, subfamily E, member 1 pseudog
NG_004308 Homo sapiens olfactory receptor, family 10, subfamily J, member 8 pseudog
NG_004309 Homo sapiens olfactory receptor, family 7, subfamily E, member 140 pseudo
NG_004310 Homo sapiens olfactory receptor, family 2, subfamily Q, member 1 pseudog
NG_004311 Homo sapiens olfactory receptor, family 52, subfamily B, member 3 pseudog
NG_004312 Homo sapiens olfactory receptor, family 8, subfamily G, member 3 pseudog
NG_004313 Homo sapiens olfactory receptor, family 7, subfamily A, member 18 pseudog
NG_004314 Homo sapiens olfactory receptor, family 7, subfamily A, member 130 pseudo
NG_004315 Homo sapiens olfactory receptor, family 7, subfamily E, member 161 pseudo
NG_004317 Homo sapiens olfactory receptor, family 55, subfamily B, member 1 pseudog
NG_004318 Homo sapiens olfactory receptor, family 51, subfamily R, member 1 pseudog
NG_004319 Homo sapiens olfactory receptor, family 52, subfamily K, member 3 pseudog
NG_004320 Homo sapiens olfactory receptor, family 51, subfamily A, member 9 pseudog
NG_004321 Homo sapiens olfactory receptor, family 51, subfamily F, member 5 pseudog
NG_004322 Homo sapiens olfactory receptor, family 51, subfamily C, member 4 pseudog
NG_004323 Homo sapiens olfactory receptor, family 51, subfamily F, member 3 pseudog
NG_004324 Homo sapiens olfactory receptor, family 51, subfamily F, member 6 pseudog
NG_004325 Homo sapiens olfactory receptor, family 51, subfamily A, member 6 pseudog
NG_004326 Homo sapiens olfactory receptor, family 56, subfamily B, member 2 pseudog
NG_004327 Homo sapiens olfactory receptor, family 4, subfamily A, member 40 pseudog
NG_004328 Homo sapiens olfactory receptor, family 4, subfamily A, member 43 pseudog
NG_004329 Homo sapiens olfactory receptor, family 4, subfamily A, member 6 pseudog
NG_004330 Homo sapiens olfactory receptor, family 4, subfamily A, member 2 pseudog
NG_004331 Homo sapiens olfactory receptor, family 4, subfamily A, member 4 pseudog
NG_004332 Homo sapiens olfactory receptor, family 4, subfamily A, member 11 pseudog
NG_004333 Homo sapiens olfactory receptor, family 4, subfamily A, member 9 pseudog
NG_004334 Homo sapiens olfactory receptor, family 4, subfamily A, member 10 pseudog
NG_004335 Homo sapiens olfactory receptor, family 4, subfamily A, member 17 pseudog
NG_004336 Homo sapiens olfactory receptor, family 7, subfamily E, member 145 pseudo
NG_004337 Homo sapiens olfactory receptor, family 2, subfamily AT, member 2 pseudog
NG_004338 Homo sapiens olfactory receptor, family 2, subfamily AT, member 1 pseudog
NG_004339 Homo sapiens olfactory receptor, family 10, subfamily N, member 1 pseudog
NG_004340 Homo sapiens olfactory receptor, family 8, subfamily F, member 1 pseudog
NG_004341 Homo sapiens olfactory receptor, family 8, subfamily A, member 2 pseudog
NG_004342 Homo sapiens olfactory receptor, family 8, subfamily B, member 10 pseudog
NG_004343 Homo sapiens olfactory receptor, family 5, subfamily BS, member 1 pseudog
NG_004344 Homo sapiens olfactory receptor, family 10, subfamily U, member 1 pseudog
NG_004345 Homo sapiens olfactory receptor, family 6, subfamily C, member 5 pseudog
NG_004346 Homo sapiens olfactory receptor, family 6, subfamily C, member 7 pseudog
NG_004347 Homo sapiens olfactory receptor, family 6, subfamily C, member 71 pseudog
NG_004348 Homo sapiens olfactory receptor, family 6, subfamily U, member 2 pseudog
NG_004349 Homo sapiens olfactory receptor, family 4, subfamily Q, member 2 pseudog
NG_004350 Homo sapiens olfactory receptor, family 4, subfamily U, member 1 pseudog
NG_004351 Homo sapiens olfactory receptor, family 4, subfamily T, member 1 pseudog
NG_004352 Homo sapiens olfactory receptor, family 11, subfamily G, member 1 pseudog
NG_004353 Homo sapiens olfactory receptor, family 11, subfamily H, member 5 pseudog
NG_004354 Homo sapiens olfactory receptor, family 11, subfamily H, member 7 pseudog
NG_004355 Homo sapiens olfactory receptor, family 4, subfamily N, member 3 pseudog
NG_004356 Homo sapiens olfactory receptor, family 4, subfamily F, member 14 pseudog
NG_004357 Homo sapiens olfactory receptor, family 4, subfamily F, member 13 pseudog
NG_004358 Homo sapiens olfactory receptor, family 4, subfamily F, member 28 pseudog
NG_004359 Homo sapiens olfactory receptor, family 4, subfamily F, member 8 pseudog
NG_004360 Homo sapiens olfactory receptor, family 7, subfamily E, member 18 pseudog

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NG_004361 Homo sapiens olfactory receptor, family 7, subfamily A, member 1 pseudoge
 NG_004362 Homo sapiens olfactory receptor, family 10, subfamily R, member 3 pseudog
 NG_004363 Homo sapiens olfactory receptor, family 6, subfamily K, member 5 pseudoge
 NG_004364 Homo sapiens olfactory receptor, family 10, subfamily J, member 2 pseudog
 NG_004365 Homo sapiens olfactory receptor, family 10, subfamily J, member 7 pseudog
 NG_004366 Homo sapiens olfactory receptor, family 10, subfamily J, member 9 pseudog
 NG_004367 Homo sapiens olfactory receptor, family 10, subfamily J, member 4 pseudog
 NG_004368 Homo sapiens olfactory receptor, family 7, subfamily E, member 23 pseudog
 NG_004369 Homo sapiens olfactory receptor, family 5, subfamily S, member 1 pseudoge
 NG_004370 Homo sapiens olfactory receptor, family 7, subfamily E, member 55 pseudog
 NG_004371 Homo sapiens olfactory receptor, family 7, subfamily E, member 35 pseudog
 NG_004372 Homo sapiens olfactory receptor, family 9, subfamily A, member 3 pseudoge
 NG_004373 Homo sapiens olfactory receptor, family 2, subfamily R, member 1 pseudoge
 NG_004374 Homo sapiens olfactory receptor, family 10, subfamily AC, member 1 pseud
 NG_004375 Homo sapiens olfactory receptor, family 2, subfamily A, member 13 pseudog
 NG_004376 Homo sapiens olfactory receptor, family 7, subfamily E, member 158 pseudo
 NG_004377 Homo sapiens olfactory receptor, family 13, subfamily E, member 1 pseudog
 NG_004378 Homo sapiens olfactory receptor, family 13, subfamily C, member 6 pseudog
 NG_004379 Homo sapiens olfactory receptor, family 2, subfamily S, member 1 pseudoge
 NG_004380 Homo sapiens olfactory receptor, family 13, subfamily D, member 2 pseudog
 NG_004381 Homo sapiens olfactory receptor, family 11, subfamily N, member 1 pseudog
 NG_004382 Homo sapiens olfactory receptor, family 3, subfamily B, member 1 pseudoge
 NG_004383 Homo sapiens spondyloepiphyseal dysplasia, late, pseudogene (LOC392597)
 NG_004384 Homo sapiens olfactory receptor, family 7, subfamily E, member 117 pseudo
 NG_004385 Homo sapiens olfactory receptor, family 7, subfamily E, member 102 pseudo
 NG_004386 Homo sapiens olfactory receptor, family 7, subfamily E, member 109 pseudo
 NG_004387 Homo sapiens olfactory receptor, family 51, subfamily A, member 8 pseudog
 NG_004388 Homo sapiens olfactory receptor, family 51, subfamily H, member 1 pseudog
 NG_004389 Homo sapiens olfactory receptor, family 51, subfamily H, member 2 pseudog
 NG_004390 Homo sapiens olfactory receptor, family 56, subfamily B, member 2 pseudog
 NG_004391 Homo sapiens olfactory receptor, family 5, subfamily BK, member 1 pseudog
 NG_004392 Homo sapiens olfactory receptor, family 11, subfamily K, member 2 pseudog
 NG_004393 Homo sapiens olfactory receptor, family 10, subfamily J, member 6 pseudog
 NG_004394 Homo sapiens olfactory receptor, family 7, subfamily E, member 46 pseudog
 NG_004395 Homo sapiens olfactory receptor, family 7, subfamily A, member 121 pseudo
 NG_004396 Homo sapiens YTH domain family 1 pseudogene (YTHDF1P) on chromosom
 NG_004397 Homo sapiens olfactory receptor, family 10, subfamily AH, member 1 pseud
 NG_004398 Homo sapiens olfactory receptor, family 7, subfamily E, member 59 pseudog
 NG_004399 Homo sapiens olfactory receptor, family 7, subfamily E, member 160 pseudo
 NG_004400 Homo sapiens olfactory receptor, family 13, subfamily D, member 3 pseudog
 NG_004401 Homo sapiens olfactory receptor, family 10, subfamily AE, member 3 pseudo
 NG_004402 Homo sapiens olfactory receptor, family 10, subfamily AK, member 1 pseudo
 NG_004403 Homo sapiens olfactory receptor, family 11, subfamily J, member 5 pseudog
 NG_004404 Homo sapiens olfactory receptor, family 11, subfamily Q, member 1 pseudog
 NG_004405 Homo sapiens olfactory receptor, family 13, subfamily Z, member 1 pseudog
 NG_004406 Homo sapiens olfactory receptor, family 13, subfamily Z, member 2 pseudog
 NG_004407 Homo sapiens olfactory receptor, family 1, subfamily M, member 4 pseudoge
 NG_004408 Homo sapiens olfactory receptor, family 2, subfamily A, member 41 pseudog
 NG_004409 Homo sapiens olfactory receptor, family 2, subfamily AO, member 1 pseudog
 NG_004410 Homo sapiens olfactory receptor, family 2, subfamily BH, member 1 pseudog
 NG_004411 Homo sapiens olfactory receptor, family 2, subfamily T, member 32 pseudog
 NG_004412 Homo sapiens olfactory receptor, family 2, subfamily X, member 1 pseudoge
 NG_004413 Homo sapiens olfactory receptor, family 4, subfamily A, member 41 pseudog
 NG_004414 Homo sapiens olfactory receptor, family 4, subfamily A, member 42 pseudog
 NG_004415 Homo sapiens olfactory receptor, family 4, subfamily A, member 44 pseudog
 NG_004416 Homo sapiens olfactory receptor, family 4, subfamily A, member 45 pseudog
 NG_004417 Homo sapiens olfactory receptor, family 4, subfamily A, member 46 pseudog

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NG_004418 Homo sapiens olfactory receptor, family 4, subfamily A, member 48 pseudog
NG_004419 Homo sapiens olfactory receptor, family 4, subfamily A, member 49 pseudog
NG_004420 Homo sapiens olfactory receptor, family 4, subfamily A, member 50 pseudog
NG_004421 Homo sapiens olfactory receptor, family 4, subfamily C, member 48 pseudog
NG_004422 Homo sapiens olfactory receptor, family 4, subfamily C, member 49 pseudog
NG_004423 Homo sapiens olfactory receptor, family 4, subfamily G, member 11 pseudog
NG_004424 Homo sapiens olfactory receptor, family 51, subfamily AB, member 1 pseudo
NG_004425 Homo sapiens olfactory receptor, family 51, subfamily B, member 8 pseudog
NG_004426 Homo sapiens olfactory receptor, family 5, subfamily AC, member 4 pseudog
NG_004427 Homo sapiens olfactory receptor, family 5, subfamily AC, member 1 pseudog
NG_004428 Homo sapiens olfactory receptor, family 5, subfamily J, member 7 pseudoge
NG_004429 Homo sapiens olfactory receptor, family 6, subfamily C, member 64 pseudog
NG_004430 Homo sapiens olfactory receptor, family 6, subfamily C, member 66 pseudog
NG_004431 Homo sapiens olfactory receptor, family 6, subfamily C, member 69 pseudog
NG_004432 Homo sapiens olfactory receptor, family 6, subfamily C, member 72 pseudog
NG_004433 Homo sapiens olfactory receptor, family 6, subfamily C, member 73 pseudog
NG_004434 Homo sapiens olfactory receptor, family 6, subfamily R, member 2 pseudoge
NG_004435 Homo sapiens olfactory receptor, family 7, subfamily E, member 155 pseudo
NG_004436 Homo sapiens olfactory receptor, family 7, subfamily E, member 159 pseudo
NG_004437 Homo sapiens olfactory receptor, family 7, subfamily G, member 15 pseudog
NG_004438 Homo sapiens olfactory receptor, family 7, subfamily H, member 2 pseudoge
NG_004439 Homo sapiens olfactory receptor, family 8, subfamily A, member 3 pseudoge
NG_004440 Homo sapiens olfactory receptor, family 8, subfamily X, member 1 pseudoge
NG_004441 Homo sapiens olfactory receptor, family 4, subfamily C, member 17 pseudog
NG_004443 Homo sapiens spondyloepiphyseal dysplasia, late, pseudogene (LOC41475)
NG_004444 Homo sapiens spondyloepiphyseal dysplasia, late, pseudogene (LOC41475;
NG_004445 Homo sapiens spondyloepiphyseal dysplasia, late, pseudogene (LOC41475;
NG_004446 Homo sapiens spondyloepiphyseal dysplasia, late, pseudogene 4 (SEDLF4)
NG_004625 Homo sapiens olfactory receptor, family 2, subfamily U, member 1 pseudoge
NG_004626 Homo sapiens olfactory receptor, family 13, subfamily I, member 1 pseudoge
NG_004627 Homo sapiens olfactory receptor, family 1, subfamily AB, member 1 pseudog
NG_004628 Homo sapiens olfactory receptor, family 8, subfamily G, member 7 pseudoge
NG_004629 Homo sapiens olfactory receptor, family 7, subfamily E, member 96 pseudog
NG_004630 Homo sapiens olfactory receptor, family 1, subfamily X, member 1 pseudoge
NG_004631 Homo sapiens olfactory receptor, family 1, subfamily X, member 5 pseudoge
NG_004632 Homo sapiens olfactory receptor, family 13, subfamily K, member 1 pseudog
NG_004633 Homo sapiens olfactory receptor, family 7, subfamily E, member 157 pseudo
NG_004634 Homo sapiens olfactory receptor, family 7, subfamily E, member 154 pseudo
NG_004635 Homo sapiens olfactory receptor, family 8, subfamily I, member 4 pseudoger
NG_004638 Homo sapiens glutathione S-transferase pi pseudogene (GSTPP) on chromo
NG_004639 Homo sapiens actin, gamma pseudogene (LOC414754) on chromosome Y
NG_004652 Homo sapiens olfactory receptor, family 2, subfamily AJ, member 1 (OR2A1
NG_004656 Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, ga
NG_004658 Homo sapiens MHC class III complement gene cluster, bimodular haplotype
NG_004662 Homo sapiens PTPN13-like, Y-linked pseudogene (LOC442865) on chromo
NG_004663 Homo sapiens PTPN13-like, Y-linked pseudogene (LOC442866) on chromo
NG_004666 Homo sapiens olfactory receptor, family 7, subfamily E, member 31 pseudog
NG_004670 Homo sapiens serine/threonine kinase 22A (spermiogenesis associated) (ST
NT_004321 Homo sapiens chromosome 1 genomic contig
NT_004350 Homo sapiens chromosome 1 genomic contig
NT_004433 Homo sapiens chromosome 1 genomic contig
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NT_004487 Homo sapiens chromosome 1 genomic contig
NT_004511 Homo sapiens chromosome 1 genomic contig
NT_004538 Homo sapiens chromosome 1 genomic contig
NT_004547 Homo sapiens chromosome 1 genomic contig
NT_004559 Homo sapiens chromosome 1 genomic contig

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NT_004610 Homo sapiens chromosome 1 genomic contig
NT_004658 Homo sapiens chromosome 1 genomic contig
NT_004671 Homo sapiens chromosome 1 genomic contig
NT_004686 Homo sapiens chromosome 1 genomic contig
NT_004754 Homo sapiens chromosome 1 genomic contig
NT_004836 Homo sapiens chromosome 1 genomic contig
NT_004873 Homo sapiens chromosome 1 genomic contig
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NT_006576 Homo sapiens chromosome 5 genomic contig
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NT_008984 Homo sapiens chromosome 11 genomic contig
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NT_009759 Homo sapiens chromosome 12 genomic contig
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NT_009952 Homo sapiens chromosome 13 genomic contig
NT_010194 Homo sapiens chromosome 15 genomic contig
NT_010274 Homo sapiens chromosome 15 genomic contig
NT_010280 Homo sapiens chromosome 15 genomic contig

NT_010393 Homo sapiens chromosome 16 genomic contig
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NT_019501 Homo sapiens chromosome 9 genomic contig
NT_019546 Homo sapiens chromosome 12 genomic contig
NT_019609 Homo sapiens chromosome 16 genomic contig
NT_019686 Homo sapiens chromosome X genomic contig
NT_021877 Homo sapiens chromosome 1 genomic contig

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NT_021937 Homo sapiens chromosome 1 genomic contig
NT_021973 Homo sapiens chromosome 1 genomic contig
NT_022052 Homo sapiens chromosome 1 genomic contig
NT_022071 Homo sapiens chromosome 1 genomic contig
NT_022135 Homo sapiens chromosome 2 genomic contig
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NT_022171 Homo sapiens chromosome 2 genomic contig
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NT_026446 Homo sapiens chromosome 15 genomic contig
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NT_026970 Homo sapiens chromosome 2 genomic contig

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NT_027140 Homo sapiens chromosome 13 genomic contig
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NT_028251 Homo sapiens chromosome 8 genomic contig
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NT_028405 Homo sapiens chromosome X genomic contig
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NT_029419 Homo sapiens chromosome 12 genomic contig
NT_029490 Homo sapiens chromosome 21 genomic contig
NT_029860 Homo sapiens chromosome 1 genomic contig
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NT_030059 Homo sapiens chromosome 10 genomic contig
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NT_030872 Homo sapiens chromosome 22 genomic contig
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gi|29793857|ref|NT_078079.1|Hs9_78148|29793857|

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g|42655573|ref|NT_086326.2|ENr114|42655573|
g|42655593|ref|NT_086327.3|ENr115|42655593|
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g|42655521|ref|NT_086342.2|ENr223|42655521|
g|42633584|ref|NT_086343.1|ENr224|42633584|
g|42655549|ref|NT_086344.3|ENr231|42655549|
g|42655523|ref|NT_086345.2|ENr232|42655523|
g|42655550|ref|NT_086346.3|ENr233|42655550|
g|42655525|ref|NT_086347.2|ENr234|42655525|

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Table II

Non-limiting examples of Stabilization Chemistries for chemically modified siNA constructs

Chemistry	pyrimidine	Purine	cap	p=S	Strand
"Stab 00"	Ribo	Ribo	TT at 3'-ends		S/AS
"Stab 1"	Ribo	Ribo	-	5 at 5'-end 1 at 3'-end	S/AS
"Stab 2"	Ribo	Ribo	-	All linkages	Usually AS
"Stab 3"	2'-fluoro	Ribo	-	4 at 5'-end 4 at 3'-end	Usually S
"Stab 4"	2'-fluoro	Ribo	5' and 3'-ends	-	Usually S
"Stab 5"	2'-fluoro	Ribo	-	1 at 3'-end	Usually AS
"Stab 6"	2'-O-Methyl	Ribo	5' and 3'-ends	-	Usually S
"Stab 7"	2'-fluoro	2'-deoxy	5' and 3'-ends	-	Usually S
"Stab 8"	2'-fluoro	2'-O-Methyl	-	1 at 3'-end	S/AS
"Stab 9"	Ribo	Ribo	5' and 3'-ends	-	Usually S
"Stab 10"	Ribo	Ribo	-	1 at 3'-end	Usually AS
"Stab 11"	2'-fluoro	2'-deoxy	-	1 at 3'-end	Usually AS
"Stab 12"	2'-fluoro	LNA	5' and 3'-ends		Usually S
"Stab 13"	2'-fluoro	LNA		1 at 3'-end	Usually AS
"Stab 14"	2'-fluoro	2'-deoxy		2 at 5'-end 1 at 3'-end	Usually AS
"Stab 15"	2'-deoxy	2'-deoxy		2 at 5'-end 1 at 3'-end	Usually AS
"Stab 16"	Ribo	2'-O-Methyl	5' and 3'-ends		Usually S
"Stab 17"	2'-O-Methyl	2'-O-Methyl	5' and 3'-ends		Usually S
"Stab 18"	2'-fluoro	2'-O-Methyl	5' and 3'-ends		Usually S
"Stab 19"	2'-fluoro	2'-O-Methyl	3'-end		S/AS
"Stab 20"	2'-fluoro	2'-deoxy	3'-end		Usually AS
"Stab 21"	2'-fluoro	Ribo	3'-end		Usually AS
"Stab 22"	Ribo	Ribo	3'-end		Usually AS
" <u>Stab 23</u> "	<u>2'-fluoro*</u>	<u>2'-deoxy*</u>	<u>5' and 3'-ends</u>		<u>Usually S</u>
" <u>Stab 24</u> "	<u>2'-fluoro*</u>	<u>2'-O-Methyl*</u>	<u>=</u>	<u>1 at 3'-end</u>	<u>S/AS</u>

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<u>"Stab 25"</u>	<u>2'-fluoro*</u>	<u>2'-O-Methyl*</u>	=	<u>1 at 3'-end</u>	<u>S/AS</u>
<u>"Stab 26"</u>	<u>2'-fluoro*</u>	<u>2'-O-Methyl*</u>	=		<u>S/AS</u>
<u>"Stab 27"</u>	<u>2'-fluoro*</u>	<u>2'-O-Methyl*</u>	3'-end		<u>S/AS</u>
<u>"Stab 28"</u>	<u>2'-fluoro*</u>	<u>2'-O-Methyl*</u>	3'-end		<u>S/AS</u>
<u>"Stab 29"</u>	<u>2'-fluoro*</u>	<u>2'-O-Methyl*</u>		<u>1 at 3'-end</u>	<u>S/AS</u>
<u>"Stab 30"</u>	<u>2'-fluoro*</u>	<u>2'-O-Methyl*</u>			<u>S/AS</u>
<u>"Stab 31"</u>	<u>2'-fluoro*</u>	<u>2'-O-Methyl*</u>	3'-end		<u>S/AS</u>
<u>"Stab 32"</u>	<u>2'-fluoro</u>	<u>2'-O-Methyl</u>			<u>S/AS</u>

CAP = any terminal cap, see for example **Figure 10**.

All Stab 00-32 chemistries can comprise 3'-terminal thymidine (TT) residues

- 5 All Stab 00-32 chemistries typically comprise about 21 nucleotides, but can vary as described herein.

S = sense strand

AS = antisense strand

*Stab 23 has a single ribonucleotide adjacent to 3'-CAP

*Stab 24 and Stab 28 have a single ribonucleotide at 5'-terminus

- 10 *Stab 25, Stab 26, and Stab 27 have three ribonucleotides at 5'-terminus

*Stab 29, Stab 30, and Stab 31, any purine at first three nucleotide positions from 5'-terminus are ribonucleotides

p = phosphorothioate linkage

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Table III

A. 2.5 µmol Synthesis Cycle ABI 394 Instrument

Reagent	Equivalents	Amount	Wait Time* DNA	Wait Time* 2'-O-methyl	Wait Time*RNA
Phosphoramidites	6.5	163 µL	45 sec	2.5 min	7.5 min
S-Ethyl Tetrazole	23.8	238 µL	45 sec	2.5 min	7.5 min
Acetic Anhydride	100	233 µL	5 sec	5 sec	5 sec
N-Methyl Imidazole	186	233 µL	5 sec	5 sec	5 sec
TCA	176	2.3 mL	21 sec	21 sec	21 sec
Iodine	11.2	1.7 mL	45 sec	45 sec	45 sec
Beaucage	12.9	645 µL	100 sec	300 sec	300 sec
Acetonitrile	NA	6.67 mL	NA	NA	NA

B. 0.2 µmol Synthesis Cycle ABI 394 Instrument

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Reagent	Equivalents	Amount	Wait Time* DNA	Wait Time* 2'-O-methyl	Wait Time* RNA
Phosphoramidites	15	31 μ L	45 sec	233 sec	465 sec
S-Ethyl Tetrazole	38.7	31 μ L	45 sec	233 min	465 sec
Acetic Anhydride	655	124 μ L	5 sec	5 sec	5 sec
N-Methyl Imidazole	1245	124 μ L	5 sec	5 sec	5 sec
TCA	700	732 μ L	10 sec	10 sec	10 sec
Iodine	20.6	244 μ L	15 sec	15 sec	15 sec
Beaucage	7.7	232 μ L	100 sec	300 sec	300 sec
Acetonitrile	NA	2.64 mL	NA	NA	NA

C. 0.2 μ mol Synthesis Cycle 96 well Instrument

Reagent	Equivalents: DNA/ 2'-O-methyl/Ribo	Amount: DNA/2'-O- methyl/Ribo	Wait Time* DNA	Wait Time* 2'-O- methyl	Wait Time* Ribo
Phosphoramidites	22/33/66	40/80/120 μ L	60 sec	180 sec	360sec
S-Ethyl Tetrazole	70/105/210	40/80/120 μ L	60 sec	180 min	360 sec
Acetic Anhydride	265/265/265	50/50/50 μ L	10 sec	10 sec	10 sec
N-Methyl Imidazole	502/502/502	50/50/50 μ L	10 sec	10 sec	10 sec
TCA	238/475/475	250/500/500 μ L	15 sec	15 sec	15 sec
Iodine	6.8/5.8/6.8	80/80/80 μ L	30 sec	30 sec	30 sec
Beaucage	34/51/51	80/120/120	100 sec	200 sec	200 sec
Acetonitrile	NA	1150/1150/1150 μ L	NA	NA	NA

- Wait time does not include contact time during delivery.
- Tandem synthesis utilizes double coupling of linker molecule

CLAIMS

What we claim is:

1. A chemically synthesized double stranded short interfering nucleic acid (siNA) molecule that directs cleavage of an expressed pseudogene target RNA via RNA interference (RNAi), wherein:
 - a) each strand of said siNA molecule is about 18 to about 23 nucleotides in length; and
 - b) one strand of said siNA molecule comprises nucleotide sequence having sufficient complementarity to said expressed pseudogene target RNA for the siNA molecule to direct cleavage of the expressed pseudogene target RNA via RNA interference.
2. The siNA molecule of claim 1, wherein said siNA molecule comprises no ribonucleotides.
3. The siNA molecule of claim 1, wherein said siNA molecule comprises one or more ribonucleotides.
4. The siNA molecule of claim 1, wherein one strand of said double-stranded siNA molecule comprises a nucleotide sequence that is complementary to a nucleotide sequence of the expressed pseudogene target RNA or a portion thereof, and wherein a second strand of said double-stranded siNA molecule comprises a nucleotide sequence substantially similar to the nucleotide sequence or a portion thereof of the expressed pseudogene target RNA.
5. The siNA molecule of claim 4, wherein each strand of the siNA molecule comprises about 18 to about 23 nucleotides, and wherein each strand comprises at least about 19 nucleotides that are complementary to the nucleotides of the other strand.
6. The siNA molecule of claim 1, wherein said siNA molecule comprises an antisense region comprising a nucleotide sequence that is complementary to a nucleotide sequence of the expressed pseudogene target RNA or a portion thereof, and wherein said siNA further comprises a sense region, wherein said sense region comprises a nucleotide sequence substantially similar to the nucleotide sequence of the expressed pseudogene target RNA or a portion thereof.

7. The siNA molecule of claim 6, wherein said antisense region and said sense region comprise about 18 to about 23 nucleotides, and wherein said antisense region comprises at least about 18 nucleotides that are complementary to nucleotides of the sense region.
8. The siNA molecule of claim 1, wherein said siNA molecule comprises a sense region and an antisense region, and wherein said antisense region comprises a nucleotide sequence that is complementary to a nucleotide sequence the expressed pseudogene target RNA or a portion thereof, and said sense region comprises a nucleotide sequence that is complementary to said antisense region.
9. The siNA molecule of claim 6, wherein said siNA molecule is assembled from two separate oligonucleotide fragments wherein one fragment comprises the sense region and a second fragment comprises the antisense region of said siNA molecule.
10. The siNA molecule of claim 6, wherein said sense region is connected to the antisense region via a linker molecule.
11. The siNA molecule of claim 10, wherein said linker molecule is a polynucleotide linker.
12. The siNA molecule of claim 10, wherein said linker molecule is a non-nucleotide linker.
13. The siNA molecule of claim 6, wherein pyrimidine nucleotides in the sense region are 2'-O-methyl pyrimidine nucleotides.
14. The siNA molecule of claim 6, wherein purine nucleotides in the sense region are 2'-deoxy purine nucleotides.
15. The siNA molecule of claim 6, wherein pyrimidine nucleotides present in the sense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides.
16. The siNA molecule of claim 9, wherein the fragment comprising said sense region includes a terminal cap moiety at a 5'-end, a 3'-end, or both of the 5' and 3' ends of the fragment comprising said sense region.
17. The siNA molecule of claim 16, wherein said terminal cap moiety is an inverted deoxy abasic moiety.

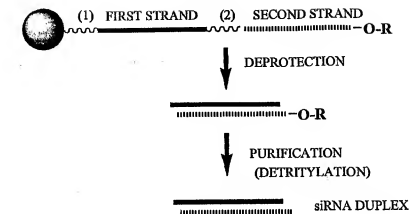
18. The siNA molecule of claim 6, wherein pyrimidine nucleotides of said antisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides.
19. The siNA molecule of claim 6, wherein purine nucleotides of said antisense region are 2'-O-methyl purine nucleotides.
20. The siNA molecule of claim 6, wherein purine nucleotides present in said antisense region comprise 2'-deoxy- purine nucleotides.
21. The siNA molecule of claim 18, wherein said antisense region comprises a phosphorothioate internucleotide linkage at the 3' end of said antisense region.
22. The siNA molecule of claim 6, wherein said antisense region comprises a glyceryl modification at a 3' end of said antisense region.
23. The siNA molecule of claim 9, wherein each of the two fragments of said siNA molecule comprise about 21 nucleotides.
24. The siNA molecule of claim 23, wherein about 19 nucleotides of each fragment of the siNA molecule are base-paired to the complementary nucleotides of the other fragment of the siNA molecule and wherein at least two 3' terminal nucleotides of each fragment of the siNA molecule are not base-paired to the nucleotides of the other fragment of the siNA molecule.
25. The siNA molecule of claim 24, wherein each of the two 3' terminal nucleotides of each fragment of the siNA molecule are 2'-deoxy-pyrimidines.
26. The siNA molecule of claim 25, wherein said 2'-deoxy-pyrimidine is 2'-deoxy-thymidine.
27. The siNA molecule of claim 23, wherein all of the about 21 nucleotides of each fragment of the siNA molecule are base-paired to the complementary nucleotides of the other fragment of the siNA molecule.
28. The siNA molecule of claim 23, wherein about 19 nucleotides of the antisense region are base-paired to the nucleotide sequence of the expressed pseudogene target RNA or a portion thereof.
29. The siNA molecule of claim 23, wherein about 21 nucleotides of the antisense region are base-paired to the nucleotide sequence of the expressed pseudogene target RNA or a portion thereof.

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30. The siNA molecule of claim 9, wherein a 5'-end of the fragment comprising said antisense region optionally includes a phosphate group.
31. A composition comprising the siNA molecule of claim 1 in an pharmaceutically acceptable carrier or diluent.
32. The siNA molecule of claim 1, wherein said expressed pseudogene target is a disease related expressed pseudogene.

Figure 1



= SOLID SUPPORT

R = TERMINAL PROTECTING GROUP

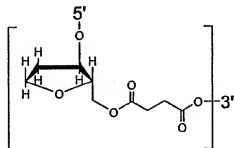
FOR EXAMPLE:
DIMETHOXYTRITYL (DMT)



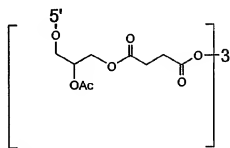
(1) = CLEAVABLE LINKER
(FOR EXAMPLE: NUCLEOTIDE SUCCINATE OR
INVERTED DEOXYABASIC SUCCINATE)



(2) = CLEAVABLE LINKER
(FOR EXAMPLE: NUCLEOTIDE SUCCINATE OR
INVERTED DEOXYABASIC SUCCINATE)



INVERTED DEOXYABASIC SUCCINATE
LINKAGE



GLYCERYL SUCCINATE LINKAGE

Figure 2

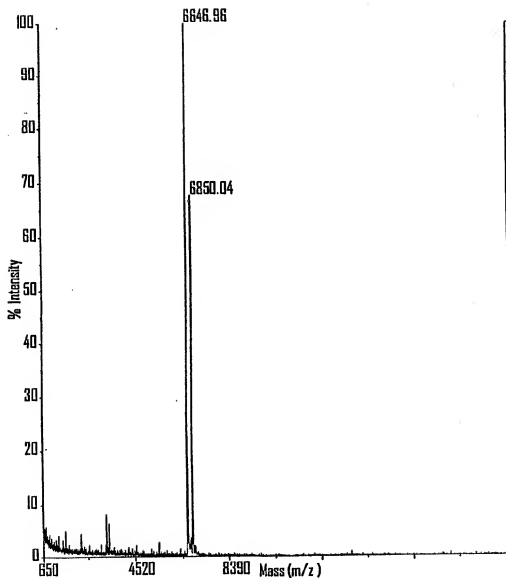


Figure 3

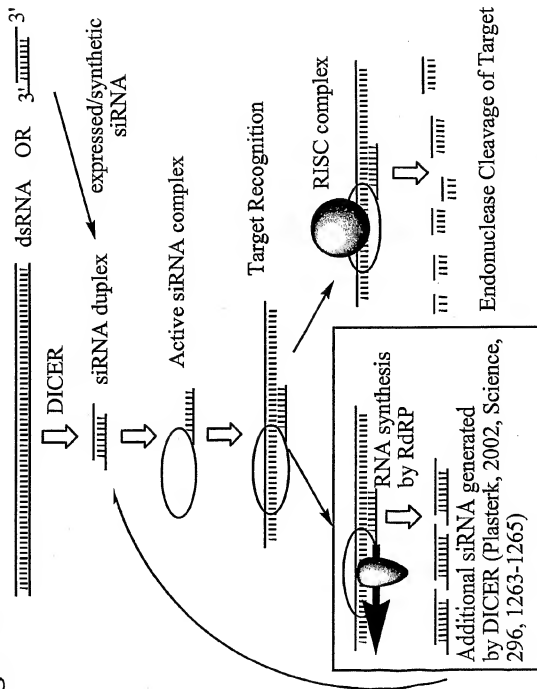
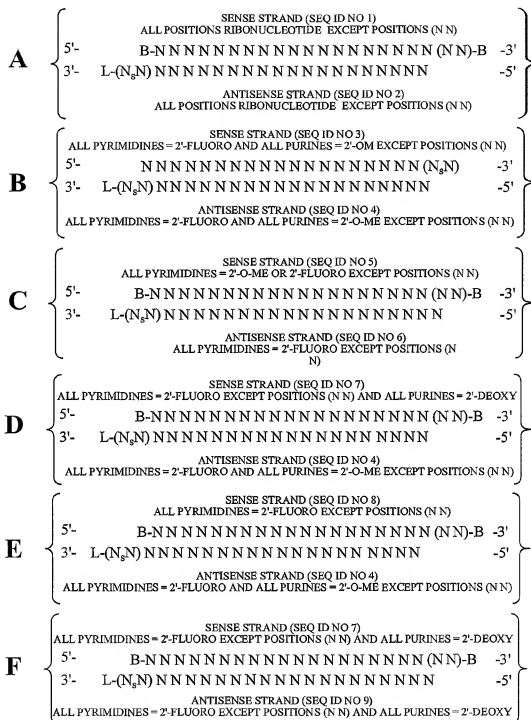
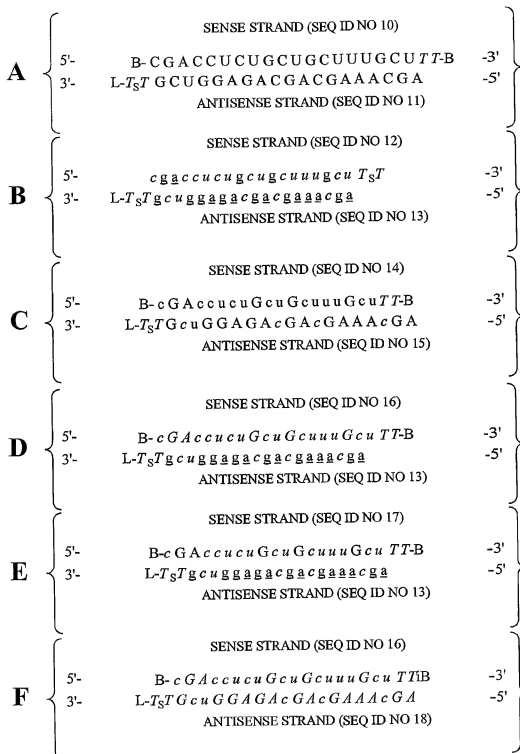


Figure 4



POSITIONS (NN) CAN COMPRISE ANY NUCLEOTIDE, SUCH AS DEOXYNUCLEOTIDES (eg. THYMIDINE) OR UNIVERSAL BASES
 B = ABASIC, INVERTED ABASIC, INVERTED NUCLEOTIDE OR OTHER TERMINAL CAP THAT IS OPTIONALLY PRESENT
 L = GLYCERYL OR B THAT IS OPTIONALLY PRESENT
 S = PHOSPHOROTHIOMATE OR PHOSPHORODITHIOATE that is optionally absent

Figure 5



lower case = 2'-O-Methyl or 2'-deoxy-2'-fluoro

italic lower case = 2'-deoxy-2'-fluoro

underline = 2'-O-methyl

ITALIC UPPER CASE = DEOXY

B = ABASIC, INVERTED ABASIC, INVERTED NUCLEOTIDE
OR OTHER TERMINAL CAP THAT IS OPTIONALLY PRESENT

L = GLYCERYL MOIETY or IB OPTIONALLY PRESENT

S = PHOSPHOROTHIOATE OR

PHOSPHORODITHIOATE OPTIONALLY PRESENT

Figure 6

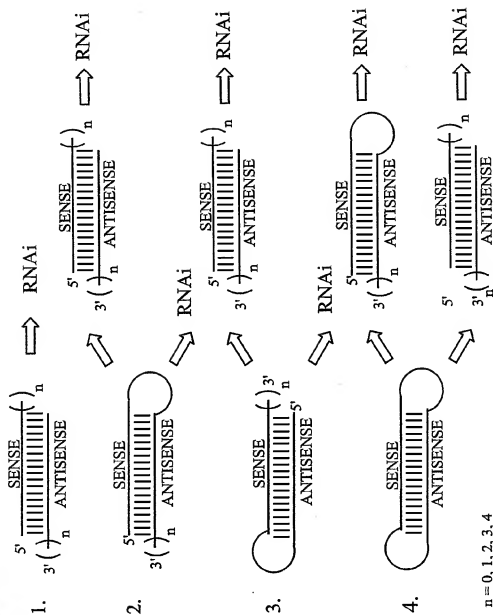
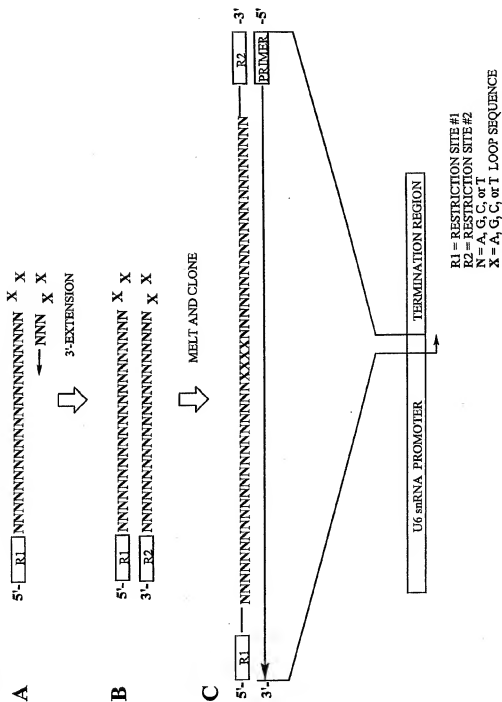


Figure 7



A

5'-[R1]NNNNNNNNNNN[R2]X X
 ↓
 [R2]X X

3'-EXTENSION

3'-EXTENSION

B

5'-R1-NNNNNNNNNNNNNNNNNNNNN-X

3'-R1-NNNNNNNNNNNNNNNNNNNNN-X

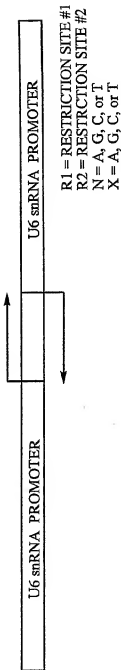
R2X
R2X



CLEAVAGE WITH RESTRICTION ENZYMES 1 AND 2

C

CLONE



R1 = RESTRICTION SITE #1
R2 = RESTRICTION SITE #2
N = A, G, C, or T
X = A, G, C, or T

Figure 9: Target site Selection using siRNA

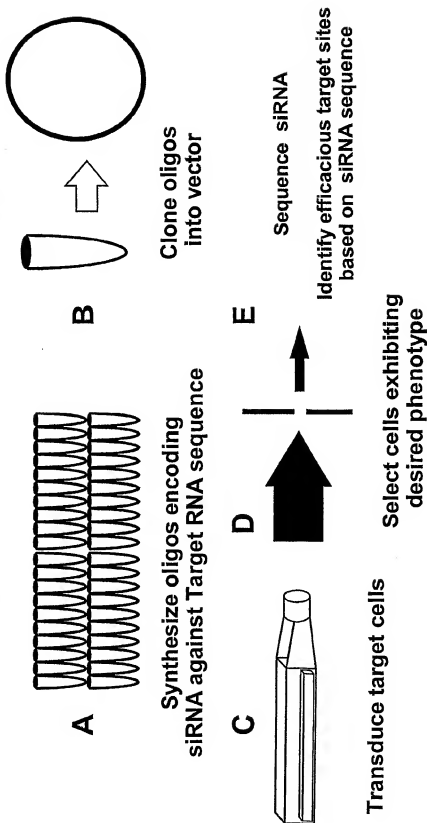
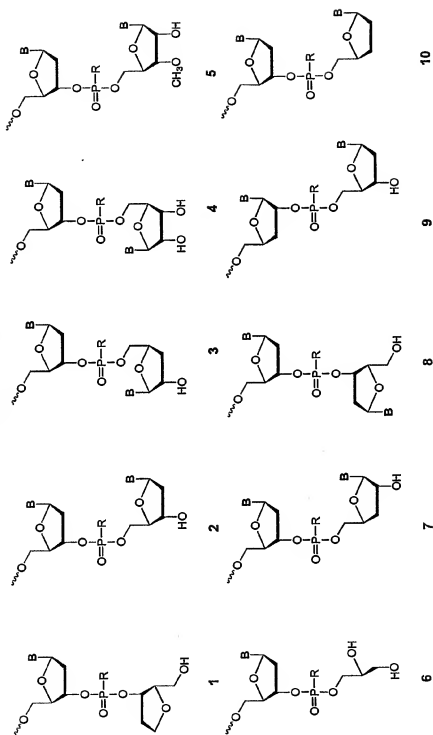


Figure 10



R = O, S, N, alkyl, substituted alkyl, O-alkyl, alkaryl, or aralkyl
 B = Independently any nucleotide base, either naturally occurring or chemically modified, or optionally H (abasic).

Figure 11: Modification Strategy

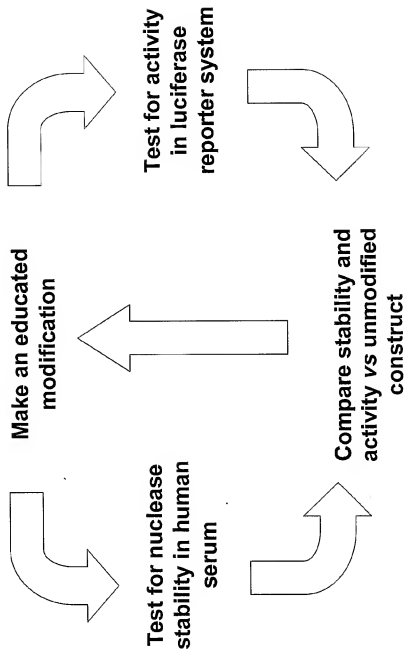
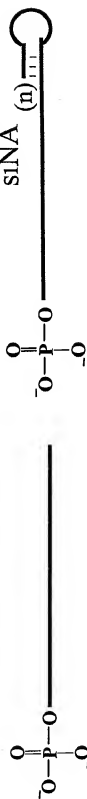


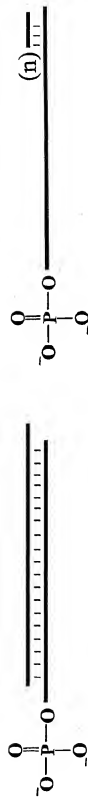
Figure 12: Phosphorylated siNA constructs



Asymmetric hairpin

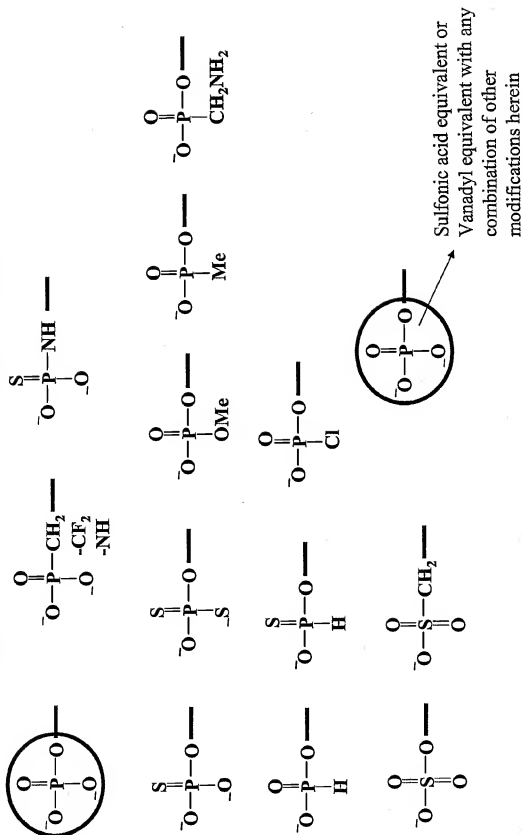
Phosphates can be modified
as described herein

Asymmetric duplex
siNA

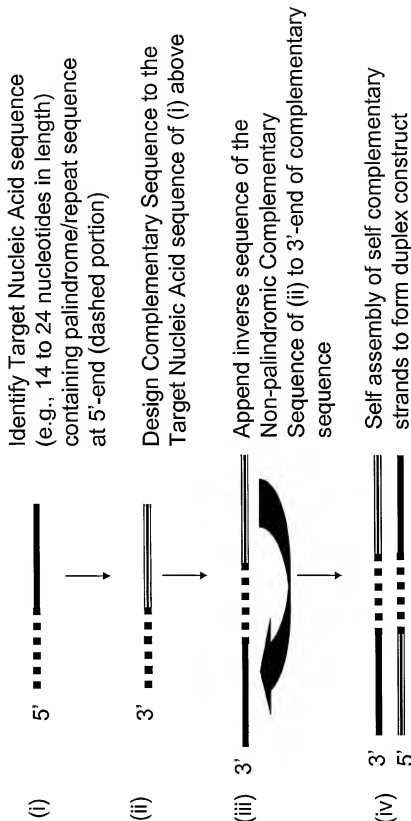


(n) = number of base
pairs (e.g. 3-18 bp)

Figure 13: 5'-phosphate modifications



**Figure 14A: Duplex forming oligonucleotide constructs that utilize
Palindromic or repeat sequences**



Identify Target Nucleic Acid sequence (e.g., 14 to 24 nucleotides in length) containing palindromic/repeat sequence at 5'-end (dashed portion)

Design Complementary Sequence to the Target Nucleic Acid sequence of (i) above

Append inverse sequence of the Non-palindromic Complementary Sequence of (ii) to 3'-end of complementary sequence

Self assembly of self complementary strands to form duplex construct (blunt ends)

SEQ ID NO: 19
5' AUAAU CUAUUUCG 3'

SEQ ID NO: 20
3' UAUAUA GAUAAAGC 5'

SEQ ID NO: 21
3' GCUUUAUC UAUAUA GAUAAAGC 5'

SEQ ID NO: 21
3' GCUUUAUC UAUAUA GAUAAAGC CGAAAUAG AUAUAU CUAUUUUCG 5'

Figure 14C: Example of a duplex forming oligonucleotide sequence that utilizes a palindrome or repeat sequence, self assembly

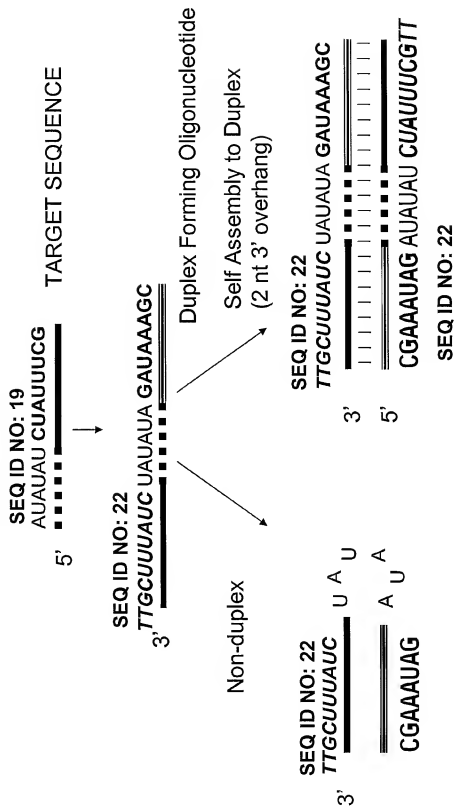


Figure 14D: Example of a duplex forming oligonucleotide sequence that utilizes a palindrome or repeat sequence, self assembly and inhibition of Target Sequence Expression

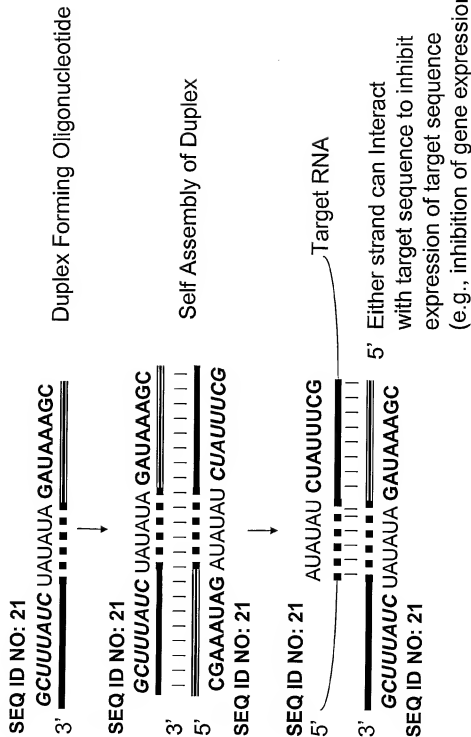


Figure 15: Duplex forming oligonucleotide constructs that utilize artificial palindrome or repeat sequences

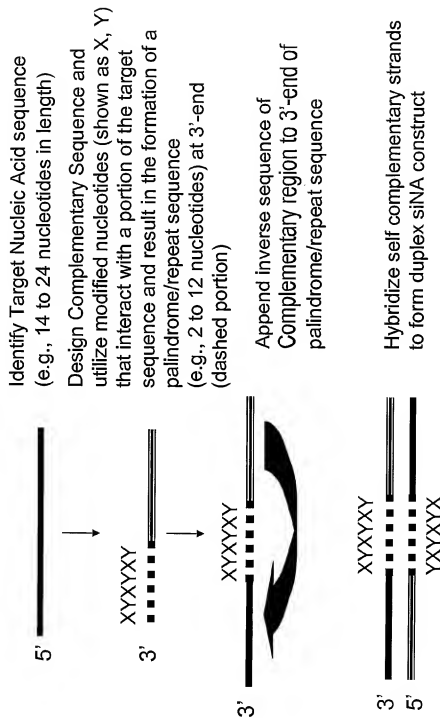


Figure 16: Examples of double stranded multifunctional siRNA constructs with distinct complementary regions

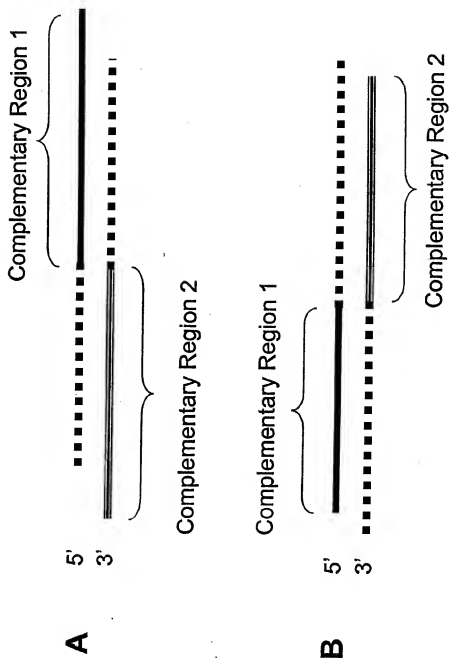


Figure 17: Examples of hairpin multifunctional siRNA constructs with distinct complementary regions

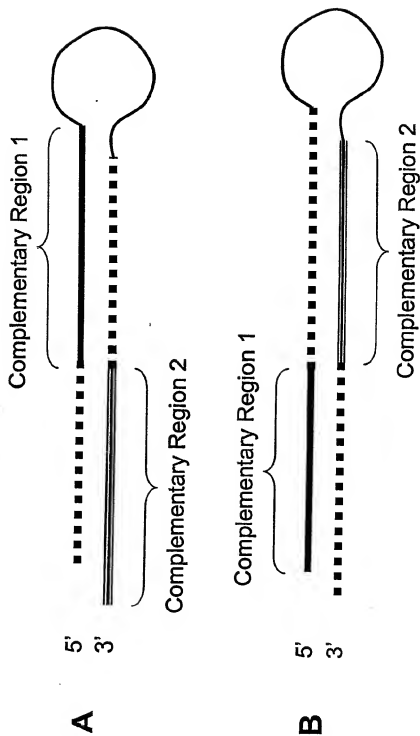


Figure 18: Examples of double stranded multifunctional siNA constructs with distinct complementary regions and a self complementary/palindrome region

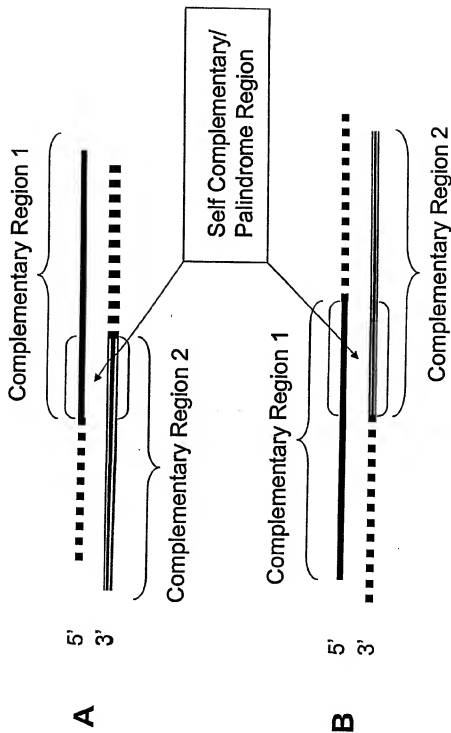
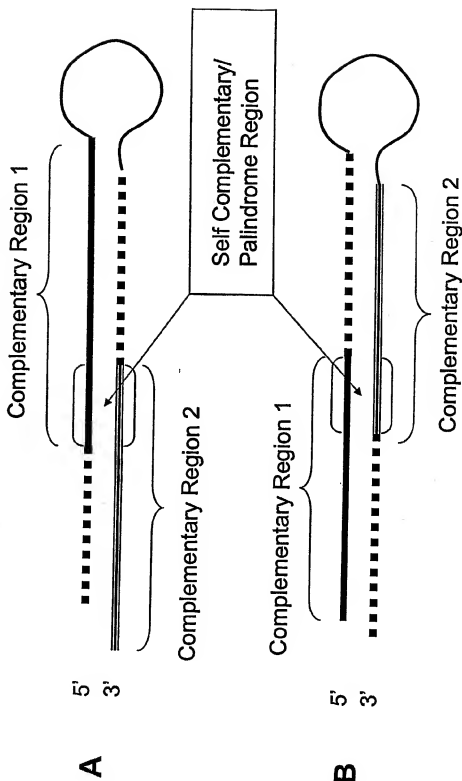


Figure 19: Examples of hairpin multifunctional siNA constructs with distinct complementary regions and a self complementary/palindrome region



**Figure 20: Example of multifunctional siRNA targeting two
Separate Target nucleic acid sequences**

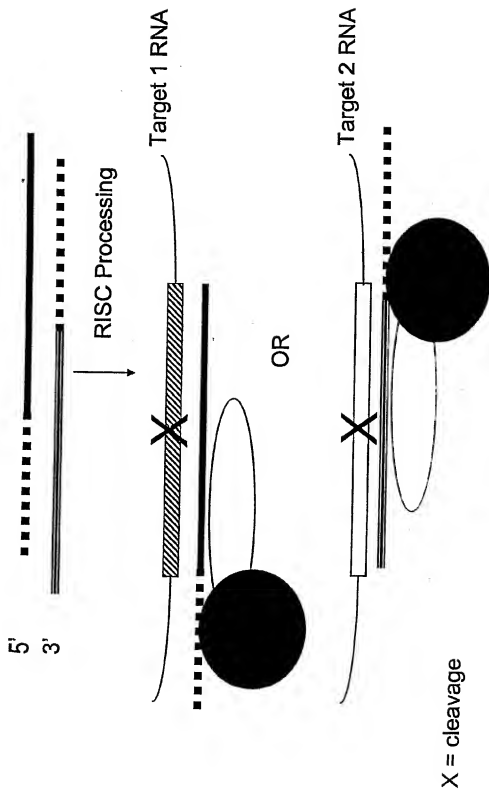


Figure 21: Example of multifunctional siRNA targeting two regions within the same target nucleic acid sequence

